GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 2, 2005, 18:23:01 ; Search time 149 Seconds
(without alignments)
785.623 Million cell updates/sec

Title: US-10-613-413B-8
Perfect score: 1717
Sequence: 1 MTPSPLLLLLLPPLLLGAFP.....VLPTGDVWSRPDGSYLNKPL 324

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1608061 seqs, 361289386 residues

Total number of hits satisfying chosen parameters: 1608061

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_AA:*
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20: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1707	99.4	357	15	US-10-613-413A-49	Sequence 49, Appl
2	1707	99.4	472	9	US-09-815-108-5	Sequence 5, Appli
3	1707	99.4	472	14	US-10-229-584-5	Sequence 5, Appli
4	1707	99.4	504	9	US-09-758-386-2	Sequence 2, Appli
5	1707	99.4	504	9	US-09-815-108-8	Sequence 8, Appli
6	1707	99.4	504	9	US-09-815-108-15	Sequence 15, Appl
7	1707	99.4	504	9	US-09-815-108-17	Sequence 17, Appl
8	1707	99.4	504	9	US-09-815-108-19	Sequence 19, Appl
9	1707	99.4	504	9	US-09-989-722-119	Sequence 119, App
10	1707	99.4	504	9	US-09-989-723-119	Sequence 119, App
11	1707	99.4	504	9	US-09-989-279-119	Sequence 119, App
12	1707	99.4	504	9	US-09-989-727-119	Sequence 119, App
13	1707	99.4	504	9	US-09-989-731-119	Sequence 119, App
14	1707	99.4	504	9	US-09-989-732-119	Sequence 119, App
15	1707	99.4	504	9	US-09-991-073-119	Sequence 119, App
16	1707	99.4	504	9	US-09-990-442-119	Sequence 119, App
17	1707	99.4	504	9	US-09-991-163-119	Sequence 119, App
18	1707	99.4	504	9	US-09-993-604-119	Sequence 119, App
19	1707	99.4	504	9	US-09-990-456-119	Sequence 119, App
20	1707	99.4	504	9	US-09-989-721-119	Sequence 119, App
21	1707	99.4	504	9	US-09-992-598-119	Sequence 119, App
22	1707	99.4	504	9	US-09-989-293A-119	Sequence 119, App
23	1707	99.4	504	9	US-09-989-735-119	Sequence 119, App
24	1707	99.4	504	9	US-09-990-444-119	Sequence 119, App
25	1707	99.4	504	9	US-09-991-181-119	Sequence 119, App
26	1707	99.4	504	9	US-09-989-730-119	Sequence 119, App
27	1707	99.4	504	9	US-09-990-436-119	Sequence 119, App
28	1707	99.4	504	9	US-09-993-687-119	Sequence 119, App
29	1707	99.4	504	10	US-09-989-734-119	Sequence 119, App
30	1707	99.4	504	10	US-09-997-653-119	Sequence 119, App
31	1707	99.4	504	10	US-09-989-724-119	Sequence 119, App
32	1707	99.4	504	10	US-09-989-728-119	Sequence 119, App
33	1707	99.4	504	10	US-09-990-441-119	Sequence 119, App
34	1707	99.4	504	10	US-09-993-667-119	Sequence 119, App
35	1707	99.4	504	10	US-09-997-428-119	Sequence 119, App
36	1707	99.4	504	10	US-09-997-666-119	Sequence 119, App
37	1707	99.4	504	10	US-09-990-438-119	Sequence 119, App
38	1707	99.4	504	10	US-09-990-562-119	Sequence 119, App
39	1707	99.4	504	10	US-09-796-753-94	Sequence 94, Appl
40	1707	99.4	504	10	US-09-796-753-108	Sequence 108, App
41	1707	99.4	504	10	US-09-990-711-119	Sequence 119, App
42	1707	99.4	504	10	US-09-989-726-119	Sequence 119, App
43	1707	99.4	504	10	US-09-998-156-119	Sequence 119, App
44	1707	99.4	504	10	US-09-990-437-119	Sequence 119, App
45	1707	99.4	504	10	US-09-991-157-119	Sequence 119, App

ALIGNMENTS

RESULT 1
US-10-613-413A-49
; Sequence 49, Application US/10613413A
; Publication No. US20040058849A1
; GENERAL INFORMATION:
; APPLICANT: Sleeman, Matthew
; APPLICANT: Murison, Greg
; TITLE OF INVENTION: Fibroblast Growth Factor Receptors and Methods for Their Use
; FILE REFERENCE: 11000.1037c5
; CURRENT APPLICATION NUMBER: US/10/613,413A
; CURRENT FILING DATE: 2003-07-03
; PRIOR APPLICATION NUMBER: U.S. 09/823,038
; PRIOR FILING DATE: 2001-03-28
; PRIOR APPLICATION NUMBER: U.S. 09/383,586
; PRIOR FILING DATE: 1999-08-26
; PRIOR APPLICATION NUMBER: U.S. 09/276,268
; PRIOR FILING DATE: 1999-03-25
; PRIOR APPLICATION NUMBER: PCT/NZ00/00015
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: U.S. 60/221,216
; PRIOR FILING DATE: 2000-07-25
; PRIOR APPLICATION NUMBER: U.S. 10/157,444
; PRIOR FILING DATE: 2000-05-28
; PRIOR APPLICATION NUMBER: PCT/NZ03/00105
; PRIOR FILING DATE: 2003-05-27
; NUMBER OF SEQ ID NOS: 145
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 49
; LENGTH: 357
; TYPE: PRT
; ORGANISM: Human
US-10-613-413A-49

Query Match 99.4%; Score 1707; DB 15; Length 357;
Best Local Similarity 99.7%; Pred. No. 4.1e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 2
US-09-815-108-5
; Sequence 5, Application US/09815108
; Patent No. US20020009776A1
; GENERAL INFORMATION:
; APPLICANT: Saris, Christiaan M.
; APPLICANT: Sharon, Mu X.
; APPLICANT: Xia, Min
; APPLICANT: Boone, Thomas Charles
; APPLICANT: Covey, Todd
; TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
; TITLE OF INVENTION: Uses Thereof
; FILE REFERENCE: 99-513-A
; CURRENT APPLICATION NUMBER: US/09/815,108
; CURRENT FILING DATE: 2001-03-22
; PRIOR APPLICATION NUMBER: 60/191,379
; PRIOR FILING DATE: 2000-03-22
; NUMBER OF SEQ ID NOS: 22
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 5
; LENGTH: 472
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-815-108-5

Query Match 99.4%; Score 1707; DB 9; Length 472;
Best Local Similarity 99.7%; Pred. No. 5.5e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

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RESULT 3
US-10-229-584-5
; Sequence 5, Application US/10229584
; Publication No. US20030087384A1
; GENERAL INFORMATION:
; APPLICANT: Saris, Christiaan M.
; APPLICANT: Sharon, Mu X.
; APPLICANT: Xia, Min
; TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
; TITLE OF INVENTION: Uses Thereof
; FILE REFERENCE: 99-513-F
; CURRENT APPLICATION NUMBER: US/10/229,584
; CURRENT FILING DATE: 2002-08-28
; PRIOR APPLICATION NUMBER: 09/815,108
; PRIOR FILING DATE: 2001-03-22
; PRIOR APPLICATION NUMBER: 60/191,379
; PRIOR FILING DATE: 2000-03-22
; NUMBER OF SEQ ID NOS: 22
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 5
; LENGTH: 472
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-229-584-5

Query Match 99.4%; Score 1707; DB 14; Length 472;
Best Local Similarity 99.7%; Pred. No. 5.5e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 4
US-09-758-386-2
; Sequence 2, Application US/09758386
; Patent No. US20010016335A1
; GENERAL INFORMATION:
; APPLICANT: Human Genome Sciences, In. et al.
; TITLE OF INVENTION: Fibroblast Growth Factor Receptor-5
; FILE REFERENCE: PF486PCT
; CURRENT APPLICATION NUMBER: US/09/758,386
; CURRENT FILING DATE: 2001-01-12
; PRIOR APPLICATION NUMBER: 09/293,182
; PRIOR FILING DATE: 1999-04-16

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;  NUMBER OF SEQ ID NOS: 15
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
;   LENGTH: 504
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-758-386-2

  Query Match             99.4%;  Score 1707;  DB 9;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 5.9e-108;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

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Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 5
US-09-815-108-8
; Sequence 8, Application US/09815108
; Patent No. US20020009776A1
; GENERAL INFORMATION:
;  APPLICANT: Saris, Christiaan M.
;  APPLICANT:  Sharon, Mu X.
;  APPLICANT:  Xia, Min
;  APPLICANT:  Boone, Thomas Charles
;  APPLICANT:  Covey, Todd
;  TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
;  TITLE OF INVENTION:  Uses Thereof
;  FILE REFERENCE: 99-513-A
;  CURRENT APPLICATION NUMBER: US/09/815,108
;  CURRENT FILING DATE:  2001-03-22
;  PRIOR APPLICATION NUMBER: 60/191,379
;  PRIOR FILING DATE: 2000-03-22
;  NUMBER OF SEQ ID NOS: 22
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 8
;   LENGTH: 504
;   TYPE: PRT
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Description of Artificial Sequence: virtual human
;   OTHER INFORMATION: FGFR-L amino acid sequence comprising residues
;   OTHER INFORMATION: 1-472 of SEQ ID NO: 5 and residues 473-504 of
;   OTHER INFORMATION: GenBank accession no. AJ277437
US-09-815-108-8

  Query Match             99.4%;  Score 1707;  DB 9;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 5.9e-108;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 6
US-09-815-108-15
; Sequence 15, Application US/09815108
; Patent No. US20020009776A1
; GENERAL INFORMATION:
;  APPLICANT: Saris, Christiaan M.
;  APPLICANT:  Sharon, Mu X.
;  APPLICANT:  Xia, Min
;  APPLICANT:  Boone, Thomas Charles
;  APPLICANT:  Covey, Todd
;  TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
;  TITLE OF INVENTION:  Uses Thereof
;  FILE REFERENCE: 99-513-A
;  CURRENT APPLICATION NUMBER: US/09/815,108
;  CURRENT FILING DATE:  2001-03-22
;  PRIOR APPLICATION NUMBER: 60/191,379
;  PRIOR FILING DATE: 2000-03-22
;  NUMBER OF SEQ ID NOS: 22
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 15
;   LENGTH: 504
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-815-108-15

  Query Match             99.4%;  Score 1707;  DB 9;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 5.9e-108;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 7
US-09-815-108-17
; Sequence 17, Application US/09815108
; Patent No. US20020009776A1
; GENERAL INFORMATION:
;  APPLICANT: Saris, Christiaan M.
;  APPLICANT:  Sharon, Mu X.
;  APPLICANT:  Xia, Min
;  APPLICANT:  Boone, Thomas Charles
;  APPLICANT:  Covey, Todd
;  TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
;  TITLE OF INVENTION:  Uses Thereof
;  FILE REFERENCE: 99-513-A
;  CURRENT APPLICATION NUMBER: US/09/815,108
;  CURRENT FILING DATE:  2001-03-22
;  PRIOR APPLICATION NUMBER: 60/191,379
;  PRIOR FILING DATE: 2000-03-22
;  NUMBER OF SEQ ID NOS: 22
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 17
;   LENGTH: 504
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-815-108-17

  Query Match             99.4%;  Score 1707;  DB 9;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 5.9e-108;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 8
US-09-815-108-19
; Sequence 19, Application US/09815108
; Patent No. US20020009776A1
; GENERAL INFORMATION:
;  APPLICANT: Saris, Christiaan M.
;  APPLICANT:  Sharon, Mu X.
;  APPLICANT:  Xia, Min
;  APPLICANT:  Boone, Thomas Charles
;  APPLICANT:  Covey, Todd
;  TITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and
;  TITLE OF INVENTION:  Uses Thereof
;  FILE REFERENCE: 99-513-A
;  CURRENT APPLICATION NUMBER: US/09/815,108
;  CURRENT FILING DATE:  2001-03-22
;  PRIOR APPLICATION NUMBER: 60/191,379
;  PRIOR FILING DATE: 2000-03-22
;  NUMBER OF SEQ ID NOS: 22
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 19
;   LENGTH: 504
;   TYPE: PRT
;   ORGANISM: Homo sapiens
US-09-815-108-19

  Query Match             99.4%;  Score 1707;  DB 9;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 5.9e-108;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 9
US-09-989-722-119
; Sequence 119, Application US/09989722
; Patent No. US20020072067A1
; GENERAL INFORMATION:
;  APPLICANT: Ashkenazi,Avi J.
;  APPLICANT:  Baker,Kevin P.
;  APPLICANT:  Botstein,David
;  APPLICANT:  Desnoyers,Luc
;  APPLICANT:  Eaton,Dan L.
;  APPLICANT:  Ferrara,Napoleone
;  APPLICANT:  Fong,Sherman
;  APPLICANT:  Gerber,Hanspeter
;  APPLICANT:  Gerritsen,Mary E.
;  APPLICANT:  Goddard,Audrey
;  APPLICANT:  Godowski,Paul J.
;  APPLICANT:  Grimaldi,J.Christopher
;  APPLICANT:  Gurney,Austin L.
;  APPLICANT:  Kljavin,Ivar J.
;  APPLICANT:  Napier,Mary A.
;  APPLICANT:  Pan,James
;  APPLICANT:  Paoni,Nicholas F.
;  APPLICANT:  Roy, Margaret Ann
;  APPLICANT:  Stewart, Timothy A.
;  APPLICANT:  Tumas, Daniel
;  APPLICANT:  Watanabe, Colin K.
;  APPLICANT:  Williams, P. Mickey
;  APPLICANT:  Wood, William I.
;  APPLICANT:  Zhang, Zemin
;  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
;  TITLE OF INVENTION:  Acids Encoding the Same
;  FILE REFERENCE: P2730P1C63
;  CURRENT APPLICATION NUMBER: US/09/989,722
;  CURRENT FILING DATE:  2001-11-19
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; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
; PRIOR FILING DATE: 1997-11-13
; PRIOR APPLICATION NUMBER: 60/066770
; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
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; PRIOR FILING DATE: 1998-06-26
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; PRIOR FILING DATE: 1998-07-01
; PRIOR APPLICATION NUMBER: 60/091478
; PRIOR FILING DATE: 1998-07-02
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; PRIOR FILING DATE: 1998-07-02
; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 10
US-09-989-723-119
; Sequence 119, Application US/09989723
; Patent No. US20020072092A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David
; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
; APPLICANT: Fong,Sherman
; APPLICANT: Gerber,Hanspeter
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Grimaldi,J.Christopher
; APPLICANT: Gurney,Austin L.
; APPLICANT: Kljavin,Ivar J.
; APPLICANT: Napier,Mary A.
; APPLICANT: Pan,James
; APPLICANT: Paoni,Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C62
; CURRENT APPLICATION NUMBER: US/09/989,723
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
; PRIOR APPLICATION NUMBER: 60/065186
; PRIOR FILING DATE: 1997-11-12
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; PRIOR FILING DATE: 1997-11-24
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; PRIOR APPLICATION NUMBER: 60/091978
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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RESULT 11
US-09-989-279-119
; Sequence 119, Application US/09989279
; Patent No. US20020072496A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David

; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
; APPLICANT: Fong,Sherman
; APPLICANT: Gerber,Hanspeter
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Grimaldi,J.Christopher
; APPLICANT: Gurney,Austin L.
; APPLICANT: Kljavin,Ivar J.
; APPLICANT: Napier,Mary A.
; APPLICANT: Pan,James
; APPLICANT: Paoni,Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C56
; CURRENT APPLICATION NUMBER: US/09/989,279
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
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; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
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RESULT 12
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; Sequence 119, Application US/09989727
; Patent No. US20020072497A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David
; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
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; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C65
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; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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RESULT 13
US-09-989-731-119
; Sequence 119, Application US/09989731
; Patent No. US20020103125A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David
; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
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; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C70
; CURRENT APPLICATION NUMBER: US/09/989,731
; CURRENT FILING DATE: 2001-11-20
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
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; PRIOR APPLICATION NUMBER: 60/091982
; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

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Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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RESULT 14
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; Sequence 119, Application US/09989732
; Patent No. US20020123463A1
; GENERAL INFORMATION:
; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David
; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
; APPLICANT: Fong,Sherman
; APPLICANT: Gerber,Hanspeter
; APPLICANT: Gerritsen,Mary E.
; APPLICANT: Goddard,Audrey
; APPLICANT: Godowski,Paul J.
; APPLICANT: Grimaldi,J.Christopher
; APPLICANT: Gurney,Austin L.
; APPLICANT: Kljavin,Ivar J.
; APPLICANT: Napier,Mary A.
; APPLICANT: Pan,James
; APPLICANT: Paoni,Nicholas F.
; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C57
; CURRENT APPLICATION NUMBER: US/09/989,732
; CURRENT FILING DATE: 2001-11-19
; PRIOR APPLICATION NUMBER: 60/049787
; PRIOR FILING DATE: 1997-06-16
; PRIOR APPLICATION NUMBER: 60/062250
; PRIOR FILING DATE: 1997-10-17
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; PRIOR FILING DATE: 1997-11-12
; PRIOR APPLICATION NUMBER: 60/065311
; PRIOR FILING DATE: 1997-11-13

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; PRIOR FILING DATE: 1997-11-24
; PRIOR APPLICATION NUMBER: 60/075945
; PRIOR FILING DATE: 1998-02-25
; PRIOR APPLICATION NUMBER: 60/078910
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; PRIOR FILING DATE: 1998-07-07
; PRIOR APPLICATION NUMBER: 60/092182
; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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RESULT 15
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; APPLICANT: Ashkenazi,Avi J.
; APPLICANT: Baker,Kevin P.
; APPLICANT: Botstein,David
; APPLICANT: Desnoyers,Luc
; APPLICANT: Eaton,Dan L.
; APPLICANT: Ferrara,Napoleone
; APPLICANT: Fong,Sherman
; APPLICANT: Gerber,Hanspeter
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; APPLICANT: Roy, Margaret Ann
; APPLICANT: Stewart, Timothy A.
; APPLICANT: Tumas, Daniel
; APPLICANT: Watanabe, Colin K.
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William I.
; APPLICANT: Zhang, Zemin
; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
; TITLE OF INVENTION: Acids Encoding the Same
; FILE REFERENCE: P2730P1C15
; CURRENT APPLICATION NUMBER: US/09/991,073
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; PRIOR FILING DATE: 1997-06-16
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; PRIOR APPLICATION NUMBER: 60/065186
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; PRIOR FILING DATE: 1998-07-09

Query Match 99.4%; Score 1707; DB 9; Length 504;
Best Local Similarity 99.7%; Pred. No. 5.9e-108;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Search completed: February 2, 2005, 18:37:32
Job time : 156 secs

GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 2, 2005, 18:12:29 ; Search time 159 Seconds
(without alignments)
730.996 Million cell updates/sec

Title: US-10-613-413B-8
Perfect score: 1717
Sequence: 1 MTPSPLLLLLLPPLLLGAFP.....VLPTGDVWSRPDGSYLNKPL 324

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2002273 seqs, 358729299 residues

Total number of hits satisfying chosen parameters: 2002273

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_23Sep04:*
1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2002s:*
6: geneseqp2003as:*
7: geneseqp2003bs:*
8: geneseqp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1717	100.0	324	8	ADF83418	Adf83418 Human fib
2	1707	99.4	472	4	AAU09810	Aau09810 Human fib
3	1707	99.4	504	3	AAY66656	Aay66656 Membrane-
4	1707	99.4	504	3	AAY92864	Aay92864 Human fib
5	1707	99.4	504	3	AAB24066	Aab24066 Human PRO
6	1707	99.4	504	4	AAB66264	Aab66264 Human MAN
7	1707	99.4	504	4	AAB65179	Aab65179 Human PRO
8	1707	99.4	504	5	AAU77790	Aau77790 Human PRO
9	1707	99.4	504	5	ABB84237	Abb84237 Human IMX
10	1707	99.4	504	5	AAU81961	Aau81961 Human PRO
11	1707	99.4	504	6	ABU57994	Abu57994 Human PRO
12	1707	99.4	504	6	ABU59072	Abu59072 Novel hum
13	1707	99.4	504	6	ABU82584	Abu82584 Human sec
14	1707	99.4	504	6	ABU60503	Abu60503 Human sec
15	1707	99.4	504	6	ABU13885	Abu13885 Human PRO
16	1707	99.4	504	6	ABU72470	Abu72470 Novel hum
17	1707	99.4	504	6	ABU59219	Abu59219 Human sec
18	1707	99.4	504	6	ABO25916	Abo25916 Human PRO
19	1707	99.4	504	6	ABU58925	Abu58925 Human sec
20	1707	99.4	504	6	ABU92303	Abu92303 Novel hum
21	1707	99.4	504	6	ABU59368	Abu59368 Novel hum
22	1707	99.4	504	6	ABU92134	Abu92134 Novel hum
23	1707	99.4	504	6	ABU10840	Abu10840 Human PRO
24	1707	99.4	504	6	ABU81592	Abu81592 Novel hum
25	1707	99.4	504	6	ABU88531	Abu88531 Human sec
26	1707	99.4	504	6	ABO34045	Abo34045 Human PRO
27	1707	99.4	504	6	ADA37630	Ada37630 Human sec
28	1707	99.4	504	6	ADA21316	Ada21316 Human sec
29	1707	99.4	504	6	ADA10103	Ada10103 Human sec
30	1707	99.4	504	6	ADA17647	Ada17647 Human PRO
31	1707	99.4	504	6	ADA27755	Ada27755 Human sec
32	1707	99.4	504	6	ADA94335	Ada94335 Human sec
33	1707	99.4	504	6	ADA38560	Ada38560 Human sec
34	1707	99.4	504	6	ADA92681	Ada92681 Human sec
35	1707	99.4	504	7	ABO53131	Abo53131 Human sec
36	1707	99.4	504	7	ADA22242	Ada22242 Human sec
37	1707	99.4	504	7	ABO22501	Abo22501 Human sec
38	1707	99.4	504	7	ADA06408	Ada06408 Human sec
39	1707	99.4	504	7	ADA39101	Ada39101 Human sec
40	1707	99.4	504	7	ADB96127	Adb96127 Human PRO
41	1707	99.4	504	7	ADC57599	Adc57599 Human PRO
42	1707	99.4	504	7	ADC54963	Adc54963 Human PRO
43	1707	99.4	504	7	ADC11830	Adc11830 Human sec
44	1707	99.4	504	7	ADC56252	Adc56252 Human PRO
45	1707	99.4	504	7	ADC07307	Adc07307 Human sec

ALIGNMENTS

RESULT 1
ADF83418
ID ADF83418 standard; protein; 324 AA.
XX
AC ADF83418;
XX
DT 26-FEB-2004 (first entry)
XX
DE Human fibroblast growth factor receptor 5 polypeptide.
XX
KW Human; fibroblast growth factor receptor 5; receptor; FGFR5; cytostatic;
KW neuroprotective; antiinflammatory; dermatological; immunosuppressive;
KW antidiabetic; antirheumatic; antiarthritic; tuberculostatic;
KW tuberculostatic; litholytic; nephrotropic; antiarteriosclerotic;
KW vasotropic; osteopathic; gene therapy.
XX
OS Homo sapiens.
XX

FH	Key	Location/Qualifiers
FT	Peptide	1. .24
FT		/note= "Signal peptide"
FT	Domain	44. .106
FT		/note= "Immunoglobulin domain"
FT	Region	51. .54
FT		/note= "CAAX box"
FT	Region	99. .102
FT		/note= "CAAX box"
FT	Domain	145. .171
FT		/note= "CAM binding domain"
FT	Domain	154. .171
FT		/note= "Heparin binding domain"
FT	Domain	165. .228
FT		/note= "Immunoglobulin domain"
FT	Modified-site	202. .205
FT		/note= "cAMP- and cGMP-dependent protein kinase
FT		phosphorylation site"
FT	Modified-site	212. .219
FT		/note= "Tyrosine kinase phosphorylation site"
FT	Region	217. .220
FT		/note= "CAAX box"
FT	Domain	261. .324
FT		/note= "Immunoglobulin domain"
FT	Region	268. .271
FT		/note= "CAAX box"

XX
PN WO2003099839-A1.
XX
PD 04-DEC-2003.

XX
PF 27-MAY-2003; 2003WO-NZ000105.
XX
PR 28-MAY-2002; 2002US-00157444.
XX
PA (GENE-) GENESIS RES & DEV CORP LTD.
XX
PI Murison JG, Sleeman M;
XX
DR WPI; 2004-035099/03.
XX
PT Modulators of fibroblast growth factor receptor 5-gene expression or
PT polypeptide function, useful in a medicament for treating a disease
PT associated with elevated osteopontin expression e.g. cancer, multiple
PT sclerosis or diabetes.
XX
PS Disclosure; Fig 10; 95pp; English.
XX
CC The present sequence is the protein sequence of human fibroblast growth
CC factor receptor 5 (FGFR5). The invention provides murine and human FGFR5
CC polypeptides and polynucleotides, as well as modulators of FGFR5 gene
CC expression and binding molecules that specifically bind to and agonise or
CC antagonise FGFR5 polypeptide function. Modulators of FGFR5 polypeptide
CC function include antibodies, scFv and Camelidae heavy chain IgG that
CC specifically bind to FGFR5. They can be used to treat diseases associated
CC with elevated osteopontin expression such as cancer (especially breast
CC cancer, hepatocellular carcinoma and colon cancer), multiple sclerosis,
CC systemic lupus erythematosus, diabetes, rheumatoid arthritis,
CC sarcoidosis, tuberculosis, kidney stones, atherosclerosis, vasculitis,
CC nephritis, arthritis, osteoporosis and osteopetrosis.
XX
SQ Sequence 324 AA;

Query Match 100.0%; Score 1717; DB 8; Length 324;
Best Local Similarity 100.0%; Pred. No. 1.7e-118;
Matches 324; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||||
Db        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
```

RESULT 2
AAU09810
ID AAU09810 standard; protein; 472 AA.
XX
AC AAU09810;
XX
DT 27-FEB-2002 (first entry)
XX
DE Human fibroblast growth factor receptor-like protein.
XX
KW Human; fibroblast growth factor receptor-like protein; FGFR-L; anorectic;
KW haemostatic; osteopathic; cytostatic; nephrotropic; antidiabetic;
KW immunomodulator; antiinflammatory; haematopoietic disorder; osteoporosis;
KW osteogenesis imperfecta; Paget's disease; periodontal disease; cancer;
KW hypercalcaemia; acute glomerulonephritis; chronic glomerulonephritis;
KW diabetes; obesity; cachexia; transgenic animal; gene therapy.
XX
OS Homo sapiens.
XX
FH Key Location/Qualifiers
FT Peptide 1. .21
FT /note= "Signal peptide"
FT Protein 22. .472
FT /note= "Mature fibroblast growth factor receptor-like
FT protein; this sequence is specifically claimed in claim
FT 39"
FT Domain 379. .399
FT /note= "Predicted transmembrane domain"
XX
PN WO200170977-A2.
XX
PD 27-SEP-2001.
XX
PF 22-MAR-2001; 2001WO-US009073.
XX
PR 22-MAR-2000; 2000US-0191379P.
XX
PA (AMGE-) AMGEN INC.
PA (SARI/) SARIS C M.
PA (MUSX/) MU S X.
PA (XIAM/) XIA M.
PA (BOON/) BOONE T C.
PA (COVE/) COVEY T.
XX
PI Saris CM, Mu SX, Xia M, Boone TC, Covey T;
XX
DR WPI; 2001-626128/72.
DR N-PSDB; AAS14936.
XX
PT Novel nucleic acid encoding fibroblast growth factor receptor-like
PT polypeptides, useful for treating hematopoietic disorder, osteoporosis,
PT Paget's disease, glomerulonephritis, cancer, diabetes, obesity and
PT cachexia.
XX
PS Claim 13; Fig 2; 163pp; English.
XX
CC The invention relates to a novel isolated fibroblast growth factor
CC receptor-like (FGFR-L) polypeptide (I). (I) and the nucleic acid (II)
CC encoding (I) are useful for treating, preventing or ameliorating a
CC medical condition including haematopoietic disorder, osteoporosis,
CC osteogenesis imperfecta, Paget's disease, periodontal disease,
CC hypercalcaemia, acute glomerulonephritis, chronic glomerulonephritis,
CC cancer, diabetes, obesity and cachexia. (I) is also useful for
CC identifying a compound which binds to FGFR-L polypeptide, by contacting
CC (I) with a compound, determining the extent of binding of the FGFR-L
CC polypeptide to the compound, and determining the activity of the
CC polypeptide when bound to the compound. (II) is useful for modulating
CC levels of a polypeptide in an animal. A transgenic animal comprising (I)
CC is useful for determining whether a compound inhibits FGFR-L polypeptide
CC activity or FGFR-L polypeptide production, by exposing the transgenic
CC animal to the compound and measuring FGFR-L polypeptide or production in
CC the mammal. (II) is useful for mapping the locations of FGFR-L gene and
CC related genes on chromosomes, as hybridisation probes in diagnostic
CC assays to test for the presence of an FGFR-L nucleic acid molecule in
CC mammalian tissue or bodily fluid samples, in gene therapy, and as tools
CC for isolating corresponding FGFR-L polypeptide genes. (I) is useful as
CC immunogen, and for cloning FGFR-L polypeptide ligands using an expression
CC cloning strategy. The present sequence represents the amino acid sequence
CC of human fibroblast growth factor receptor-like protein as described in
CC the invention
XX
SQ Sequence 472 AA;

Query Match 99.4%; Score 1707; DB 4; Length 472;

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  Best Local Similarity   99.7%;  Pred. No. 1.4e-117;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 3
AAY66656
ID   AAY66656 standard; protein; 504 AA.
XX
AC   AAY66656;
XX
DT   05-APR-2000  (first entry)
XX
DE   Membrane-bound protein PRO943.
XX
KW   Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;
KW   pharmaceutical; receptor immunoadhesin; gene mapping.
XX
OS   Homo sapiens.
XX
PN   WO9963088-A2.
XX
PD   09-DEC-1999.
XX
PF   02-JUN-1999;   99WO-US012252.
XX
PR   02-JUN-1998;   98US-0087607P.
PR   02-JUN-1998;   98US-0087609P.
PR   02-JUN-1998;   98US-0087759P.
PR   03-JUN-1998;   98US-0087827P.
PR   04-JUN-1998;   98US-0088021P.
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PR   04-AUG-1998;   98US-0095318P.
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PR 16-SEP-1998; 98US-0100634P.
PR 12-JAN-1999; 99US-0115565P.
XX
PA (GETH) GENENTECH INC.
XX
PI Baker K, Chen J, Goddard A, Gurney AL, Smith V, Watanabe CK;
PI Wood WI, Yuan J;
XX
DR WPI; 2000-072883/06.
DR N-PSDB; AAZ64984.
XX
PT Membrane-bound proteins and related nucleotide sequences.
XX
PS Claim 12; Fig 70; 822pp; English.
XX
CC The invention provides membrane-bound PRO polypeptides and
CC polynucleotides encoding them. The PRO sequences of the invention were
CC identified based on extracellular domain homology screening. The PRO
CC sequences have homology with proteins including LDL receptors, TIE
CC ligands and various enzymes. The membrane-bound proteins and receptor
CC molecules are useful as pharmaceutical and diagnostic agents. Receptor
CC immunoadhesins, for instance, can be used as therapeutic agents to block
CC receptor-ligand interactions. The membrane-bound proteins can also be
CC employed for screening of potential peptide or small molecule inhibitors
CC of the relevant receptor/ligand interaction. The PRO encoding sequences
CC are useful as hybridization probes, in chromosome and gene mapping and in
CC the generation of antisense RNA and DNA. PRO nucleic acid sequences will
CC also be useful for the preparation of PRO polypeptides, especially by
CC recombinant techniques
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 3; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 4
AAY92864
ID AAY92864 standard; protein; 504 AA.
XX
AC AAY92864;
XX
DT 29-AUG-2000 (first entry)
XX
DE Human fibroblast growth factor receptor 5.
XX
KW FGFR-5; fibroblast growth factor receptor 5; cytostatic; anti-sclerotic;
KW immunomodulatory; gastrointestinal; virucide; anti-inflammatory;
KW anti-ischemic; anti-atherosclerosis; angiogenic; endocrine;
KW anti-diabetic; gene therapy.
XX
OS Homo sapiens.
XX

FH	Key	Location/Qualifiers
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FT	Peptide	23. .37
FT		/label= antigenic
FT	Protein	25. .504
FT		/label= mature_protein
FT	Domain	25. .117
FT		/label= extracellular
FT		/note= "immunoglobulin domain I"
FT	Peptide	39. .48
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FT	Peptide	51. .59
FT		/label= antigenic
FT	Peptide	62. .76
FT		/label= antigenic
FT	Peptide	81. .97
FT		/label= antigenic
FT	Peptide	101. .104
FT		/label= antigenic
FT	Domain	118. .143
FT		/label= acid_box_domain
FT	Peptide	119. .170
FT		/label= antigenic
FT	Domain	144. .239
FT		/label= extracellular
FT		/note= "immunoglobulin domain II"
FT	Peptide	176. .204
FT		/label= antigenic
FT	Peptide	209. .228
FT		/label= antigenic

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FT   Peptide          238. .247
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FT   Domain           240. .357
FT                    /label= extracellular
FT                    /note= "immunoglobulin domain III"
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FT   Peptide          326. .334
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XX
PN   WO200024756-A1.
XX
PD   04-MAY-2000.
XX
PF   17-JUN-1999;   99WO-US013620.
XX
PR   23-OCT-1998;   98US-0105465P.
XX
PA   (HUMA-) HUMAN GENOME SCI INC.
XX
PI   Ruben SM,  Young PE;
XX
DR   WPI; 2000-387035/33.
DR   N-PSDB; AAA28842.
XX
PT   Nucleic acids encoding fibroblast growth factor-5 useful for the
PT   prevention, diagnosis and treatment of conditions associated with tissue
PT   repair and aberrant cell functions, e.g. cell survival and proliferation.
XX
PS   Claim 11; Fig 1A-C; 182pp; English.
XX
CC   This is the fibroblast growth factor receptor protein, FGFR-5. The FGFR-5
CC   protein and DNA may be used in the prevention, treatment and diagnosis of
CC   diseases or conditions associated with inappropriate FGFR-5 expression
CC   and activity. For example, the nucleic acids (and vectors containing
CC   them) and the FGFR-5 polypeptide may be used to treat disorders
CC   associated with increased or decreased cell survival (such as cancers
CC   (e.g. leukemia, colonic cancer, testicular cancer and follicular
CC   lymphomas), autoimmune disorders (e.g. multiple sclerosis and Crohn's
CC   disease) viral infections (e.g. herpes viruses), inflammation, graft
CC   versus host disease, acute and chronic graft rejection, ischemic injuries
CC   and atherosclerosis), activation, secretion, migration, differentiation
CC   and proliferation, diseases associated with defects in wound healing,
CC   mucositosis, defects of angiogenesis, immune dysfunction, endocrine
CC   dysfunction and insulin secretion disorders. Anti-FGFR-5 antibodies may
CC   also be used as diagnostic agents for detecting the presence of FGFR-5
CC   polypeptides in samples
XX
SQ   Sequence 504 AA;
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```
  Query Match             99.4%;  Score 1707;  DB 3;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 1.5e-117;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
```

```
RESULT 5
AAB24066
ID   AAB24066 standard; protein; 504 AA.
XX
AC   AAB24066;
XX
DT   29-JAN-2001  (first entry)
XX
DE   Human PRO943 protein sequence SEQ ID NO:29.
XX
KW   Human; tumour; diagnosis; neoplastic disease; neoplastic cell growth;
KW   proliferation; tumourigenesis; identification; cancer; cytostatic;
KW   nootropic; neuroprotective; antiinflammatory; immunosuppressive;
KW   immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy;
KW   neuronal disorder; glial disorder; astrocytal disorder; angiogenic;
KW   hypothalamic disorder; glandular disorder; macrophagal disorder;
KW   epithelial disorder; stromal disorder; blastocoelic disorder;
KW   inflammatory disorder; immunologic disorder.
XX
OS   Homo sapiens.
XX
PN   WO200053755-A2.
XX
PD   14-SEP-2000.
XX
PF   06-JAN-2000; 2000WO-US000376.
XX
PR   08-MAR-1999;   99WO-US005028.
PR   02-JUN-1999;   99WO-US012252.
PR   23-JUN-1999;   99US-0141037P.
PR   07-JUL-1999;   99US-0143048P.
PR   26-JUL-1999;   99US-0145698P.
PR   30-NOV-1999;   99WO-US028313.
PR   20-DEC-1999;   99WO-US030911.
PR   05-JAN-2000; 2000WO-US000219.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Baker KP,  Goddard A,  Gurney AL,  Hillan KJ,  Roy MA;
PI   Watanabe CK,  Wood WI;
XX
DR   WPI; 2000-572270/53.
DR   N-PSDB; AAC58376.
```

XX
PT Thirty PRO polynucleotides encoding PRO polypeptides, useful in the
PT treatment, diagnosis and prevention of cancer.
XX
PS Claim 61; Fig 20; 286pp; English.
XX
CC The present invention describes an isolated antibody that binds to one of
CC the human PRO proteins designated PRO212, PRO290, PRO341, PRO535, PRO619,
CC PRO717, PRO809, PRO830, PRO848, PRO943, PRO1005, PRO1009, PRO1025,
CC PRO1030, PRO1097, PRO1107, PRO1111, PRO1153, PRO1182, PRO1184, PRO1187,
CC PRO1281, PRO23, PRO39, PRO834, PRO1317, PRO1710, PRO2094, PRO2145 OR
CC PRO2198. PRO antagonists can be used to inhibit tumour cell growth. The
CC PRO polypeptides and nucleotides are useful in the treatment, diagnosis
CC and prevention of cancer. The antibodies and other anti-tumour compounds
CC maybe used to treat various conditions, including those characterised by
CC overexpression and/or activation of the amplified PRO genes. Exemplary
CC conditions or disorders to be treated with such antibodies and other
CC compounds include benign or malignant tumours (e.g., renal, liver,
CC kidney, bladder, breast, gastric, ovarian, colorectal, prostate,
CC pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas,
CC glioblastomas, and various head and neck tumours), leukaemias and
CC lymphoid malignancies, other disorders such as neuronal, glial,
CC astrocytal, hypothalamic and other glandular, macrophagal, epithelial,
CC stromal and blastocoelic disorders, and inflammatory, angiogenic and
CC immunologic disorders. AAC58242 to AAC58366 represent PCR primers and
CC hybridisation probes used in the isolation of the human PRO sequences.
CC AAC58367 to AAC58396 and AAB24057 to AAB24089 represent human PRO
CC polynucleotide and protein sequences given in the exemplification of the
CC present invention
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 3; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
```

RESULT 6
AAB66264
ID AAB66264 standard; protein; 504 AA.
XX
AC AAB66264;
XX
DT 05-APR-2001 (first entry)
XX
DE Human MANGO 003 SEQ ID NO: 5.
XX
KW Membrane associated protein; secreted protein; human; mouse; rat;
KW INTERCEPT 340; MANGO 003; MANGO 347; TANGO 272; TANGO 295; TANGO 354;
KW TANGO 378; skeletal disorder; cardiovascular disorder; renal disorder;
KW haematopoietic disorder; neural disorder; hepatic disorder;
KW neoplastic disease.
XX
OS Homo sapiens.
XX
PN WO200100673-A1.
XX
PD 04-JAN-2001.
XX
PF 29-JUN-2000; 2000WO-US018198.
XX
PR 30-JUN-1999; 99US-00345464.
XX
PA (MILL-) MILLENNIUM PHARM INC.
XX
PI Barnes TM, Fraser CC, Wrighton N, Myers P, Busfield SJ, Sharp JD;
XX
DR WPI; 2001-050128/06.
DR N-PSDB; AAF27781.
XX
PT Isolated secreted or transmembrane proteins are used for diagnosis and
PT treatment of neoplastic and hematopoietic disorders e.g. T cell
PT disorders, cancer and tumors.
XX
PS Claim 9; Page 216-217; 294pp; English.
XX
CC The present invention provides the protein and coding sequences for a
CC number of membrane associated and secreted proteins from human, mouse and
CC rat. The proteins are designated INTERCEPT 340, MANGO 003, MANGO 347,
CC TANGO 272, TANGO 295, TANGO 254 and TANGO 378. The proteins are all
CC involved in signal transduction and the sequences can be used in the
CC treatment of cardiovascular, renal, hepatic, neural, neoplastic, skeletal
CC and haematopoietic disorders
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 4; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy.         1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
```

RESULT 7
AAB65179
ID AAB65179 standard; protein; 504 AA.
XX
AC AAB65179;
XX
DT 02-APR-2001 (first entry)

XX
DE Human PRO943 (UNQ480) protein sequence SEQ ID NO:119.
XX
KW Human; secreted and transmembrane protein; PRO; cytostatic; cell death;
KW cancer; chromosomal mapping; gene mapping; tissue typing;
KW diagnostic assay.
XX
OS Homo sapiens.
XX
PN WO200073454-A1.
XX
PD 07-DEC-2000.
XX
PF 30-MAR-2000; 2000WO-US008439.
XX
PR 02-JUN-1999; 99WO-US012252.
PR 23-JUN-1999; 99US-0141037P.
PR 07-JUL-1999; 99US-0143048P.
PR 20-JUL-1999; 99US-0144758P.
PR 26-JUL-1999; 99US-0145698P.
PR 28-JUL-1999; 99US-0146222P.
PR 17-AUG-1999; 99US-0149396P.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 08-OCT-1999; 99US-0158663P.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
PI Grimaldi CJ, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX
DR WPI; 2001-032160/04.
DR N-PSDB; AAF44130.
XX
PT PRO polynucleotides used to produce polypeptides used to target bioactive
PT molecules such as toxins, radiolabels or antibodies, to specific cells,
PT to cause targeted cell death.
XX
PS Claim 12; Fig 70; 935pp; English.
XX
CC The present invention describes human secreted and transmembrane PRO
CC proteins. The PRO proteins have cytostatic activity. The PRO proteins can
CC be used for targeted delivery of bioactive molecules, such as toxins,
CC radiolabels or antibodies, that cause cell death. PRO nucleotide
CC sequences, and their fragments, can be used as hybridisation probes, in
CC chromosomal and gene mapping, and in the generation of anti-sense RNA and
CC DNA. They may also be used to produce transgenic animals which are used
CC to develop and screen therapeutically useful reagents. The PRO nucleotide
CC and protein sequence can be used for tissue typing and in treating
CC cancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to
CC AAF44470 represent PCR primers and hybridisation probes used in the
CC isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to
CC AAB65300 represent human PRO polynucleotide and protein sequences given
CC in the exemplification of the present invention
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 4; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```
Qy    1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy   61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy  121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy  181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy  241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db  241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy  301 QKFVVLPTGDVWSRPDGSYLNKPL 324
          |||||||||||||||||||||| |
Db  301 QKFVVLPTGDVWSRPDGSYLNKLL 324
```

RESULT 8
AAU77790
ID AAU77790 standard; protein; 504 AA.
XX
AC AAU77790;
XX
DT 05-JUN-2002 (first entry)
XX
DE Human PRO943 protein.
XX
KW PRO; cancer; neoplastic; human; tumour; PRO943; PRO1250; PRO1337;
KW breast cancer; ovarian cancer; colorectal cancer; lung cancer;
KW central nervous system cancer; melanoma; leukaemia.
XX
OS Homo sapiens.
XX
PN WO200149715-A2.
XX
PD 12-JUL-2001.
XX
PF 08-NOV-2000; 2000WO-US030952.
XX
PR 06-JAN-2000; 2000WO-US000376.
PR 18-FEB-2000; 2000WO-US004342.
PR 02-MAR-2000; 2000WO-US005841.
PR 30-MAR-2000; 2000WO-US008439.
PR 28-JUL-2000; 2000WO-US020710.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi AJ, Goddard A, Gurney AL, Napier MA, Watanabe CK;
PI Wood WI;
XX
DR WPI; 2002-256031/30.
DR N-PSDB; ABK11750.
XX
PT Composition for inhibiting neoplastic cell growth or treating tumors
PT e.g.breast cancer and ovarian cancer in mammals comprising PRO943,
PT PRO1250 or PRO1337 or its agonist.
XX
PS Claim 19; Fig 2; 101pp; English.
XX
CC This sequence relates to a novel composition useful for inhibiting

```
CC   neoplastic cell growth or treating tumours in mammals. The method
CC   comprises a polypeptide PRO943, PRO1250 or PRO1337 of the invention in
CC   admixture with a carrier. The invention also comprises the nucleotide and
CC   protein sequences of the three PRO proteins. The composition of the
CC   invention is useful for inhibiting neoplastic cell growth and for
CC   treating tumours e.g. breast cancer, ovarian cancer, colorectal cancer,
CC   lung cancer, central nervous system cancer, melanoma or leukaemia in a
CC   mammal. An expression vector containing the nucleotide sequences of the
CC   invention is useful for producing the PRO proteins by recombinant
CC   techniques, these proteins may then be used to create anti-PRO antibodies
CC   which may be used for detection of PRO protein or inhibition of PRO
CC   protein activity. The present sequence represents the human PRO943
CC   protein of the invention
XX
SQ   Sequence 504 AA;

  Query Match             99.4%;  Score 1707;  DB 5;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 1.5e-117;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 9
ABB84237
ID   ABB84237 standard; protein; 504 AA.
XX
AC   ABB84237;
XX
DT   20-SEP-2002  (first entry)
XX
DE   Human IMXP-888 isoform PRO943 protein.
XX
KW   Human; IMXP-888; PRO943; fibroblast growth factor receptor; virucide;
KW   antibacterial; fungicide; cytostatic; antirheumatic; antiarthritic;
KW   antibacterial; immunosuppressive; neuroprotective; antiinflammatory;
KW   antidiabetic; dermatological; antiasthmatic; antiallergic; vasotropic;
KW   antiatherosclerotic; antipsoriatic; nootropic; infection; cancer; lupus;
KW   graft versus host disorder; inflammatory disorder; rheumatoid arthritis;
KW   septic shock; multiple sclerosis; adult respiratory distress syndrome;
KW   ARDS; pneumonia; diabetes; asthma; allergy; reperfusion injury; eczema;
KW   atherosclerosis; cardiovascular disease; psoriasis; fibrosis;
KW   sarcoidosis; Alzheimer's disease; cytokine inducer.
XX
OS   Homo sapiens.
XX
FH   Key             Location/Qualifiers
FT   Peptide         1. .17
FT                   /label= signal_peptide
FT   Region          18. .375
FT                   /note= "soluble extracellular region specifically claimed
FT                   in Claim 4"
FT   Region          23. .370
FT                   /note= "soluble extracellular region specifically claimed
FT                   in Claim 5"
FT   Region          376. .396
FT                   /note= "transmembrane region"
XX
PN   WO200255533-A2.
XX
PD   18-JUL-2002.
XX
PF   20-NOV-2001; 2001WO-US043782.
XX
PR   22-NOV-2000; 2000US-0252785P.
XX
PA   (IMMV ) IMMUNEX CORP.
XX
PI   Chipman SD,  Schooley KA,  Born TL,  Dubose RF;
XX
DR   WPI; 2002-557812/59.
XX
PT   Activating immune system in a mammal by administering to the mammal a
PT   cytokine inducer polypeptide, IMXP-888, or treating inflammatory disorder
PT   in a mammal by administering an IMXP-888 antagonist to the mammal.
XX
PS   Claim 4; Page 39-41; 41pp; English.
XX
CC   This invention describes a novel method for activating the immune system
CC   in a mammal. The method comprises administering to the mammal an
CC   effective amount of a cytokine inducer polypeptide, IMXP-888, or treating
CC   an inflammatory disorder in a mammal comprising, administering an
CC   effective amount of an IMXP-888 antagonist to the mammal. The method of
CC   the invention has virucide, antibacterial, fungicide, cytostatic,
CC   antirheumatic, antiarthritic, antibacterial, immunosuppressive,
CC   neuroprotective, antiinflammatory, antidiabetic, dermatological,
CC   antiasthmatic, antiallergic, vasotropic, antiatherosclerotic,
CC   antipsoriatic and nootropic activity. The method is useful for activating
CC   the immune system in a mammal having a condition selected from viral,
CC   bacterial or fungal infection, cancer, and graft versus host disorders,
CC   or for treating an inflammatory disorder in a mammal, preferably human.
CC   The method is useful for treating inflammatory disorders such as
CC   rheumatoid arthritis, septic shock, multiple sclerosis, adult respiratory
CC   distress syndrome (ARDS), pneumonia, diabetes, lupus, asthma and other
CC   lung conditions, allergies, reperfusion injury, atherosclerosis and other
CC   cardiovascular diseases, eczema, psoriasis, fibrosis and the range of
CC   fibrotic disorders, sarcoidosis, Alzheimer's disease, and cancer. This
CC   sequence represents member of the IMXP-888 family, PRO943, used in the
CC   method of the invention
XX
SQ   Sequence 504 AA;

  Query Match             99.4%;  Score 1707;  DB 5;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 1.5e-117;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 10
AAU81961
ID   AAU81961 standard; protein; 504 AA.
XX
AC   AAU81961;
XX
DT   09-APR-2002  (first entry)
XX
DE   Human PRO943.
XX
KW   Human; PRO; antiinflammatory; ophthalmological; vasotropic;
KW   retinal cell injury; ocular disease; retinitis pigmentosa;
KW   macular degeneration; retinal detachment; retinal tear; retinopathy;
KW   retinal degenerative disease; macular hole; degenerative myopia;
KW   acute retinal necrosis syndrome; traumatic chorioretinopathy;
KW   Purtscher's retinopathy; oedema; ischaemic condition;
KW   retinal vision occlusion; collagen vascular disease;
KW   thrombocytopaenic purpura; uveitis; retinal vasculitis; Eales disease;
KW   systemic lupus erythematosus; environmental trauma.
XX
OS   Homo sapiens.
XX
PN   WO200109327-A2.
XX
PD   08-FEB-2001.
XX
PF   28-JUL-2000; 2000WO-US020710.
XX
PR   28-JUL-1999;   99US-0146222P.
PR   13-SEP-1999;   99WO-US020944.
PR   15-SEP-1999;   99WO-US021090.
PR   29-NOV-1999;   99WO-US028214.
PR   30-NOV-1999;   99WO-US028313.
PR   01-DEC-1999;   99WO-US028301.
PR   05-JAN-2000; 2000WO-US000219.
PR   06-JAN-2000; 2000WO-US000376.
PR   11-FEB-2000; 2000WO-US003565.
PR   18-FEB-2000; 2000WO-US004341.
PR   22-FEB-2000; 2000WO-US004414.
PR   24-FEB-2000; 2000WO-US005004.
PR   02-MAR-2000; 2000WO-US005841.
PR   15-MAR-2000; 2000WO-US006884.
PR   30-MAR-2000; 2000WO-US008439.
PR   17-MAY-2000; 2000WO-US013705.
XX
PA   (GETH ) GENENTECH INC.
XX
PI   Ashkenazi AJ,  Baker KP,  Goddard A,  Godowski PJ,  Gurney AL;
PI   Kljavin IJ,  Lafleur M,  Mark MR,  Marsters SA,  Pitti RM,  Watanabe CK;
PI   Wood WI;
XX
DR   WPI; 2002-130120/17.
DR   N-PSDB; ABK28591.
XX
PT   Promoting survival of retinal cells, or delaying or preventing retinal
PT   cell injury or death, by contacting retinal cells with PRO175, 220, 216,
PT   243, 306, 346, 322, 536, 943, 840, 828, 826, 1068 or PRO1132 polypeptide.
XX
PS   Claim 44; Fig 19; 152pp; English.
XX
CC   The invention relates to promoting the survival of retinal cells, or
CC   delaying or preventing retinal cell injury or death, by contacting the
CC   retinal cells with the polypeptide such as PRO175, PRO220, PRO216,
CC   PRO243, PRO306, PRO346, PRO322, PRO536, PRO943, PRO840, PRO828, PRO826,
CC   PRO1068 or PRO1132 polypeptide. Also included are the nucleic acids
CC   encoding the PRO proteins, a vector comprising the nucleic acid, a host
CC   cell comprising the vector, and anti-PRO antibody. The PRO proteins are
CC   useful for promoting survival of retinal cells (retinal neurons such as
CC   retinal ganglion cells, displaced retinal ganglion cells, amacrine cells,
CC   displaced amacrine cells, horizontal neurons or bipolar neurons, rod
CC   photoreceptors, or supportive cells such as Muller cells or pigment
CC   epithelial cells), or delaying or preventing retinal cell injury or death
CC   caused by ocular disease (which is or is associated with retinitis
CC   pigmentosa, macular degeneration, retinal detachment, retinal tear,
CC   retinopathy, retinal degenerative disease, macular hole, degenerative
CC   myopia, acute retinal necrosis syndrome, traumatic chorioretinopathy or
CC   contusion, Purtscher's retinopathy, oedema, an ischaemic condition,
CC   central or branch retinal vision occlusion, collagen vascular disease,
CC   thrombocytopaenic purpura, uveitis, retinal vasculitis, occlusion
CC   associated with Eales disease or systemic lupus erythematosus), retinal
CC   injury or environmental trauma. The retinal cell injury or death is
CC   delayed or prevented by substantially not causing angiogenesis or
CC   mitogenesis. The present sequence represents a PRO protein
XX
SQ   Sequence 504 AA;

  Query Match             99.4%;  Score 1707;  DB 5;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 1.5e-117;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 11
ABU57994
ID   ABU57994 standard; protein; 504 AA.
XX
AC   ABU57994;
XX
DT   14-APR-2003  (first entry)
XX
DE   Human PRO polypeptide #26.
XX
KW   Human; PRO; cytostatic; tumour; cancer; breast; lung; stomach; liver;
KW   horse; cow; dog; cat; sheep; pig; goat; rabbit; ADEPT;
KW   antibody-dependent enzyme mediated prodrug therapy.
XX
OS   Homo sapiens.
XX
PN   US2003027163-A1.
XX
PD   06-FEB-2003.
XX
PF   15-NOV-2001; 2001US-00997666.
XX
PR   16-JUN-1997;   97US-0049787P.
PR   17-OCT-1997;   97US-0062250P.
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PR 05-NOV-1997; 97WO-US020069.
PR 12-NOV-1997; 97US-0065186P.
PR 13-NOV-1997; 97US-0065311P.
PR 24-NOV-1997; 97US-0066770P.
PR 25-FEB-1998; 98US-0075945P.
PR 20-MAR-1998; 98US-0078910P.
PR 28-APR-1998; 98US-0083322P.
PR 07-MAY-1998; 98US-0084600P.
PR 28-MAY-1998; 98US-0087106P.
PR 02-JUN-1998; 98US-0087607P.
PR 02-JUN-1998; 98US-0087609P.
PR 02-JUN-1998; 98US-0087759P.
PR 03-JUN-1998; 98US-0087827P.
PR 04-JUN-1998; 98US-0088021P.
PR 04-JUN-1998; 98US-0088025P.
PR 04-JUN-1998; 98US-0088026P.
PR 04-JUN-1998; 98US-0088028P.
PR 04-JUN-1998; 98US-0088029P.
PR 04-JUN-1998; 98US-0088030P.
PR 04-JUN-1998; 98US-0088033P.
PR 04-JUN-1998; 98US-0088326P.
PR 05-JUN-1998; 98US-0088167P.
PR 05-JUN-1998; 98US-0088202P.
PR 05-JUN-1998; 98US-0088212P.
PR 05-JUN-1998; 98US-0088217P.
PR 09-JUN-1998; 98US-0088655P.
PR 10-JUN-1998; 98US-0088734P.
PR 10-JUN-1998; 98US-0088738P.
PR 10-JUN-1998; 98US-0088742P.
PR 10-JUN-1998; 98US-0088810P.
PR 10-JUN-1998; 98US-0088824P.
PR 10-JUN-1998; 98US-0088826P.
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Query Match 99.4%; Score 1707; DB 6; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
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RESULT 12
ABU59072
ID ABU59072 standard; protein; 504 AA.
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AC ABU59072;
XX
DT 28-APR-2003 (first entry)
XX
DE Novel human secreted or transmembrane protein PRO943.
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KW Human; PRO; hypertrophy of neonatal heart; angiogenesis; wound healing;
KW cardiac insufficiency disorder; cancer; tumour; immune response;
KW adrenal cortical capillary endothelial growth; c-fos induction;
KW vascular endothelial growth factor inhibition; VEGF inhibition;
KW endothelial cell growth inhibitor; T-lymphocytes stimulation;
KW retinal neurons cell survival; rod photoreceptor cell survival;
KW retinal disorder; retinitis pigmentosum; kidney disorder;
KW mammalian kidney mesangial cell proliferation; Berger disease;
KW dermatitis; herpetiformis; Crohn's disease; chondrocyte proliferation;
KW chondrocyte redifferentiation; sports injury; arthritis.
XX
OS Homo sapiens.
XX
PN US2002132252-A1.
XX
PD 19-SEP-2002.
XX
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PR 16-JUN-1997; 97US-0049787P.
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XX
PA (GETH) GENENTECH INC.
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PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
PI Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX
DR WPI; 2003-247083/24.
DR N-PSDB; ABX80197.
XX
PT Novel isolated PRO polypeptides e.g., PRO826, PRO1068, PRO1184, PRO1346
PT and PRO1375, which stimulate proliferation of stimulated T-lymphocytes
PT are therapeutically useful for enhancing immune response and in cancer
PT treatments.
XX
PS Claim 12; Fig 70; 648pp; English.
XX
CC The invention describes an isolated human PRO polypeptide. The PRO
CC polypeptides are useful in detecting PRO polypeptides in a sample, in
CC linking a bioactive molecule to a cell expressing a PRO polypeptide, and
CC in modulating at least one biological activity of a cell expressing a PRO
CC polypeptide. PRO1312 stimulates hypertrophy of neonatal heart and is thus
CC useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186
CC stimulate adrenal cortical capillary endothelial growth, and PRO536,
CC PRO943, PRO828, PRO826, PRO1068 or PRO535, PRO826, PRO819, PRO1126,
CC PRO1360 and PRO1387 induce c-fos in endothelial cells, and are thus
CC useful for treating conditions or disorders where angiogenesis would be
CC beneficial, e.g. wound healing and antagonist of this polypeptide are
CC useful for treating cancerous tumours. PRO812 inhibits vascular
CC endothelial growth factor (VEGF) stimulated proliferation of endothelial
CC cells and is thus useful for inhibiting endothelial cell growth in
CC mammals which would be beneficial in inhibiting tumour growth. PRO826,
CC PRO1068, PRO1184, PRO1346 and PRO1375 stimulate proliferation of
CC stimulated T-lymphocytes and are therapeutically useful for enhancing
CC immune response. PRO828, PRO826, PRO1068 or PRO1132 enhance survival of
CC retinal neurons cells (PRO1132 is also enhances survival/proliferation of
CC rod photoreceptor cells) and therefore are useful for treating retinal
CC disorders of injuries, e.g. retinitis pigmentosum, AMD. PRO819, PRO813
CC and PRO11066 induce proliferation of mammalian kidney mesangial cells,
CC and therefore are useful for treating kidney disorders associated with
CC decreased mesangial cell function such as Berger disease or other
CC nephropathies associated with dermatitis, herpetiformis or Crohn's
CC disease. PRO1310, PRO844, PRO1312, PRO1192 and PRO1387 induce the
CC proliferation and/or redifferentiation of chondrocytes in culture and are
CC thus useful for treating sports injuries, and arthritis. This is the
CC amino acid sequence of a novel human PRO protein
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 6; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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RESULT 13
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ID ABU82584 standard; protein; 504 AA.
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AC ABU82584;
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DT 26-JUN-2003 (first entry)
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DE Human secreted/transmembrane protein PRO943.
XX
KW Human; PRO; secreted protein; transmembrane protein;
KW cardiac insufficiency disorders; angiogenesis; wound healing;
KW cancerous tumour; immune response; retinal disorder; sight loss;
KW retinitis pigmentosum; age-related macular degeneration; AMD;
KW kidney disorder; Berger disease; nephropathy; dermatitis; herpetiformis;
KW Crohn's disease; sports injury; arthritis.
XX
OS Homo sapiens.
XX
PN US2003032023-A1.
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PD 13-FEB-2003.
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PR 17-SEP-1998; 98US-0100858P.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
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PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 12-MAR-1999; 99US-0123957P.
PR 02-JUN-1999; 99WO-US012252.
PR 23-JUN-1999; 99US-0141037P.
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PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
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PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 15-MAY-2000; 2000WO-US013358.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 23-JUN-2000; 2000US-0213637P.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.

Query Match 99.4%; Score 1707; DB 6; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 14
ABU60503
ID ABU60503 standard; protein; 504 AA.
XX
AC ABU60503;
XX
DT 01-MAY-2003 (first entry)
XX
DE Human secreted/transmembrane protein, #44.
XX
KW Human; PRO; secreted; transmembrane; signal peptide; pharmaceutical;
KW diagnostic; therapeutic; gene therapy.
XX
OS Homo sapiens.
XX

PN US2002160384-A1.
XX
PD 31-OCT-2002.
XX
PF 14-NOV-2001; 2001US-00992598.
XX
PR 16-JUN-1997; 97US-0049787P.
PR 17-OCT-1997; 97US-0062250P.
PR 05-NOV-1997; 97WO-US020069.
PR 12-NOV-1997; 97US-0065186P.
PR 13-NOV-1997; 97US-0065311P.
PR 24-NOV-1997; 97US-0066770P.
PR 25-FEB-1998; 98US-0075945P.
PR 20-MAR-1998; 98US-0078910P.
PR 28-APR-1998; 98US-0083322P.
PR 07-MAY-1998; 98US-0084600P.
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PR 02-JUN-1998; 98US-0087607P.
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PR 04-JUN-1998; 98US-0088026P.
PR 04-JUN-1998; 98US-0088028P.
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PR 05-JUN-1998; 98US-0088167P.
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PR 09-JUN-1998; 98US-0088655P.
PR 10-JUN-1998; 98US-0088734P.
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PR 16-JUN-1998; 98US-0089512P.
PR 16-JUN-1998; 98US-0089514P.
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PR 17-JUN-1998; 98US-0089538P.
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PR 17-JUN-1998; 98US-0089599P.
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PR 17-JUN-1998; 98US-0089653P.
PR 18-JUN-1998; 98US-0089801P.
PR 18-JUN-1998; 98US-0089907P.
PR 18-JUN-1998; 98US-0089908P.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 02-JUN-1999; 99WO-US012252.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 05-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.

PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 15-MAY-2000; 2000WO-US013358.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 01-DEC-2000; 2000WO-US032678.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-JUN-2001; 2001WO-US017800.
PR 20-JUN-2001; 2001WO-US019692.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 28-AUG-2001; 2001US-00941992.
XX
PA (GETH) GENENTECH INC.
XX
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
PI Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX
DR WPI; 2003-288106/28.
DR N-PSDB; ABX90175.
XX
PT New transmembrane polypeptides and nucleic acids encoding the
PT polypeptides, useful in gene therapy, in chromosome identification, as
PT chromosome markers, or in generating probes.
XX
PS Claim 12; Fig 70; 650pp; English.
XX
CC The invention discloses isolated PRO secreted/transmembrane polypeptides
CC comprising a sequence without signal peptide and the nucleic acid
CC encoding them. The polypeptides can be used to raise antibodies that
CC specifically bind to the PRO polypeptide, for linking a bioactive
CC molecule to a cell expressing a PRO protein and for modulating at least
CC one biological activity of a cell. The PRO polypeptides or
CC polynucleotides are also useful in gene therapy, in chromosome
CC identification, as chromosome markers, or in generating probes. The PRO
CC polypeptides are useful as molecular markers for protein electrophoresis,
CC and the isolated nucleic acids may be used for recombinantly expressing
CC those markers. The PRO polypeptides and nucleic acids may also be used in
CC tissue typing. Anti-PRO antibodies are useful in diagnostic assays for
CC PRO, and in affinity purification of PRO from recombinant cell culture or
CC natural sources. The sequences presented in ABU60478-ABU60624 are the PRO
CC polynucleotides of the invention. Note: The sequence data for this patent
CC is also available in electronic format from USPTO at
CC seqdata.uspto.gov/sequence.html
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 6; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
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Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 15
ABU13885
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AC ABU13885;
XX
DT 26-FEB-2003 (first entry)
XX
DE Human PRO943 polypeptide.
XX
KW Human; PRO polypeptide; secreted protein; transmembrane protein;
KW genetic disorder; antibacterial; immunosuppressive.
XX
OS Homo sapiens.
XX
PN US2002103125-A1.
XX
PD 01-AUG-2002.
XX
PF 20-NOV-2001; 2001US-00989731.
XX
PR 16-JUN-1997; 97US-0049787P.
PR 17-OCT-1997; 97US-0062250P.
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PR 20-MAR-1998; 98US-0078910P.
PR 28-APR-1998; 98US-0083322P.
PR 07-MAY-1998; 98US-0084600P.
PR 28-MAY-1998; 98US-0087106P.
PR 02-JUN-1998; 98US-0087607P.
PR 02-JUN-1998; 98US-0087609P.
PR 02-JUN-1998; 98US-0087759P.
PR 03-JUN-1998; 98US-0087827P.
PR 04-JUN-1998; 98US-0088021P.
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PR 18-JUN-1998; 98US-0089801P.
PR 18-JUN-1998; 98US-0089907P.
PR 18-JUN-1998; 98US-0089908P.
PR 16-SEP-1998; 98WO-US019330.
PR 17-SEP-1998; 98WO-US019437.
PR 07-OCT-1998; 98WO-US021141.
PR 01-DEC-1998; 98WO-US025108.
PR 05-JAN-1999; 99WO-US000106.
PR 08-MAR-1999; 99WO-US005028.
PR 02-JUN-1999; 99WO-US012252.
PR 15-SEP-1999; 99WO-US021090.
PR 15-SEP-1999; 99WO-US021547.
PR 30-NOV-1999; 99WO-US028313.
PR 01-DEC-1999; 99WO-US028301.
PR 01-DEC-1999; 99WO-US028634.
PR 16-DEC-1999; 99WO-US030095.
PR 20-DEC-1999; 99WO-US030911.
PR 06-JAN-2000; 2000WO-US000219.
PR 06-JAN-2000; 2000WO-US000376.
PR 11-FEB-2000; 2000WO-US003565.
PR 18-FEB-2000; 2000WO-US004341.
PR 22-FEB-2000; 2000WO-US004414.
PR 24-FEB-2000; 2000WO-US004914.
PR 24-FEB-2000; 2000WO-US005004.
PR 02-MAR-2000; 2000WO-US005841.
PR 10-MAR-2000; 2000WO-US006319.
PR 15-MAR-2000; 2000WO-US006884.
PR 20-MAR-2000; 2000WO-US007377.
PR 30-MAR-2000; 2000WO-US008439.
PR 15-MAY-2000; 2000WO-US013358.
PR 17-MAY-2000; 2000WO-US013705.
PR 22-MAY-2000; 2000WO-US014042.
PR 30-MAY-2000; 2000WO-US014941.
PR 02-JUN-2000; 2000WO-US015264.
PR 28-JUL-2000; 2000WO-US020710.
PR 11-AUG-2000; 2000WO-US022031.
PR 23-AUG-2000; 2000WO-US023522.
PR 24-AUG-2000; 2000WO-US023328.
PR 08-NOV-2000; 2000WO-US030952.
PR 01-DEC-2000; 2000WO-US032678.
PR 28-FEB-2001; 2001WO-US006520.
PR 01-JUN-2001; 2001WO-US017800.
PR 20-JUN-2001; 2001WO-US019692.
PR 29-JUN-2001; 2001WO-US021066.
PR 09-JUL-2001; 2001WO-US021735.
PR 28-AUG-2001; 2001US-00941992.
XX
PA (GETH) GENENTECH LTD.
XX
PI Ashkenazi AJ, Baker KP, Botstein D, Desnoyers L, Eaton DL;
PI Ferrara N, Fong S, Gerber H, Gerritsen ME, Goddard A, Godowski PJ;
PI Grimaldi JC, Gurney AL, Kljavin IJ, Napier MA, Pan J, Paoni NF;
PI Roy MA, Stewart TA, Tumas D, Watanabe CK, Williams PM, Wood WI;
PI Zhang Z;
XX
DR WPI; 2003-102117/09.
DR N-PSDB; ABX64021.
XX
PT Novel secreted and transmembrane polypeptide for modulating biological
PT activity of cell expressing the polypeptide, identifying agonists or
PT antagonists of polypeptide, and as molecular weight markers.
XX
PS Claim 12; Fig 70; 649pp; English.
XX
CC The present invention relates to the isolation of novel human PRO
CC polypeptides, and the polynucleotide sequences encoding them. The PRO
CC polypeptides are secreted and transmembrane proteins. The PRO
CC polypeptides are useful for detecting other PRO polypeptides, for linking
CC bioactive molecules to cells expressing PRO polypeptides, for modulating
CC biological activities of cells expressing PRO polypeptides, and for for
CC identifying agonists or antagonists. The polynucleotide sequences
CC encoding PRO polypeptides are useful as hybridisation probes, in
CC chromosome and gene mapping, in the generation of antisense RNA and DNA,
CC in the preparation of PRO polypeptides, for generating transgenic animals
CC or knockout animals, to construct hybridisation probes for mapping the
CC gene which encodes the PRO polypeptide, and for the genetic analysis of
CC individuals with genetic disorders, in gene therapy, for chromosome
CC identification, as chromosome markers, and for generating probes for PCR,
CC Northern analysis, Southern analysis and Western analysis. ABU13860-
CC ABU14006 represent the human PRO polypeptides of the invention. Note: The
CC sequence data for this patent was obtained in electronic format directly
CC from the USPTO web site at seqdata.uspto.gov/psipsDIDEntry.html
XX
SQ Sequence 504 AA;

Query Match 99.4%; Score 1707; DB 6; Length 504;
Best Local Similarity 99.7%; Pred. No. 1.5e-117;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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Search completed: February 2, 2005, 18:22:43
Job time : 163 secs

GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 2, 2005, 18:19:11 ; Search time 24 Seconds
(without alignments)
895.293 Million cell updates/sec

Title: US-10-613-413B-8
Perfect score: 1717
Sequence: 1 MTPSPLLLLLLPPLLLGAFP.....VLPTGDVWSRPDGSYLNKPL 324

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 478139 seqs, 66318000 residues

Total number of hits satisfying chosen parameters: 478139

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
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6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1560	90.9	373	4	US-09-823-038A-60	Sequence 60, Appl
2	1560	90.9	529	3	US-09-383-586-31	Sequence 31, Appl
3	1560	90.9	529	4	US-09-823-038A-31	Sequence 31, Appl
4	1458	84.9	322	3	US-09-383-586-33	Sequence 33, Appl
5	1458	84.9	322	4	US-09-823-038A-33	Sequence 33, Appl
6	1035.5	60.3	439	3	US-09-383-586-32	Sequence 32, Appl
7	1035.5	60.3	439	4	US-09-823-038A-32	Sequence 32, Appl
8	408.5	23.8	801	3	US-09-383-630-6	Sequence 6, Appli
9	393	22.9	806	3	US-09-383-630-3	Sequence 3, Appli
10	382.5	22.3	802	3	US-09-173-151A-33	Sequence 33, Appl
11	380	22.1	816	1	US-07-640-029-1	Sequence 1, Appli
12	379.5	22.1	355	1	US-08-471-570-14	Sequence 14, Appl
13	379.5	22.1	643	1	US-08-471-570-6	Sequence 6, Appli
14	379.5	22.1	769	1	US-08-471-570-8	Sequence 8, Appli
15	375.5	21.9	821	2	US-08-451-822A-13	Sequence 13, Appl
16	375.5	21.9	821	3	US-08-323-430-13	Sequence 13, Appl
17	371.5	21.6	622	4	US-09-499-846-2	Sequence 2, Appli
18	371.5	21.6	820	1	US-07-921-807B-3	Sequence 3, Appli
19	371.5	21.6	820	1	US-08-441-944A-3	Sequence 3, Appli
20	371.5	21.6	820	3	US-08-439-992A-1	Sequence 1, Appli
21	368.5	21.5	351	5	PCT-US93-05703-2	Sequence 2, Appli
22	367.5	21.4	817	1	US-07-640-029-2	Sequence 2, Appli
23	367.5	21.4	822	1	US-07-921-807B-4	Sequence 4, Appli
24	367.5	21.4	822	1	US-08-459-296-2	Sequence 2, Appli
25	367.5	21.4	822	1	US-08-441-944A-4	Sequence 4, Appli
26	367.5	21.4	822	2	US-08-451-822A-12	Sequence 12, Appl
27	367.5	21.4	822	3	US-08-439-992A-2	Sequence 2, Appli
28	367.5	21.4	822	3	US-08-323-430-12	Sequence 12, Appl
29	366.5	21.3	822	1	US-07-997-133-1	Sequence 1, Appli
30	357.5	20.8	820	1	US-08-166-717D-6	Sequence 6, Appli
31	350	20.4	126	3	US-09-383-586-30	Sequence 30, Appl
32	350	20.4	126	4	US-09-823-038A-30	Sequence 30, Appl
33	308	17.9	300	1	US-07-640-029-5	Sequence 5, Appli
34	308	17.9	300	3	US-08-439-992A-5	Sequence 5, Appli
35	308	17.9	525	4	US-09-499-846-4	Sequence 4, Appli
36	308	17.9	525	4	US-09-499-846-8	Sequence 8, Appli
37	307	17.9	526	1	US-08-471-570-4	Sequence 4, Appli
38	307	17.9	652	1	US-08-471-570-10	Sequence 10, Appl
39	306	17.8	302	1	US-07-921-807B-7	Sequence 7, Appli
40	306	17.8	302	1	US-08-441-944A-7	Sequence 7, Appli
41	305	17.8	240	1	US-08-471-570-12	Sequence 12, Appl
42	304	17.7	302	1	US-07-640-029-6	Sequence 6, Appli
43	304	17.7	302	1	US-07-921-807B-8	Sequence 8, Appli
44	304	17.7	302	1	US-08-441-944A-8	Sequence 8, Appli
45	304	17.7	302	3	US-08-439-992A-6	Sequence 6, Appli

ALIGNMENTS

RESULT 1
US-09-823-038A-60
; Sequence 60, Application US/09823038A
; Patent No. 6797271
; GENERAL INFORMATION:
; APPLICANT: Strachan, Lorna
; APPLICANT: Sleeman, Matthew
; APPLICANT: Abernethy, Nevin
; APPLICANT: Onrust, Rene
; APPLICANT: Kumble, Anand
; APPLICANT: Murison, Greg
; TITLE OF INVENTION: Compositions Isolated From Stromal Cells
; TITLE OF INVENTION: and Methods For Their Use
; FILE REFERENCE: 11000.1037c3
; CURRENT APPLICATION NUMBER: US/09/823,038A
; CURRENT FILING DATE: 2001-07-09
; NUMBER OF SEQ ID NOS: 61
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 60
; LENGTH: 373
; TYPE: PRT
; ORGANISM: Mouse
US-09-823-038A-60

Query Match 90.9%; Score 1560; DB 4; Length 373;
Best Local Similarity 91.4%; Pred. No. 8.3e-135;
Matches 296; Conservative 9; Mismatches 15; Indels 4; Gaps 1;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    ||||| | | ||||||:||||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||:|| |||||||||||||||||||||:::|||
Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||||
Db        117 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| ||||||||||:||||||||||:|||||||||||||||
Db        177 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||:||||||||||||
Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 2
US-09-383-586-31
; Sequence 31, Application US/09383586
; Patent No. 6242419
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT: Sleeman, Matthew
;  APPLICANT: Abernethy, Nevin
;  APPLICANT: Onrust, Rene
;  APPLICANT: Kumble, Anand
;  APPLICANT: Murison, Greg
;  TITLE OF INVENTION: Compounds isolated from stromal cells
;  TITLE OF INVENTION: and methods for their use
;  FILE REFERENCE: 11000.1037c1
;  CURRENT APPLICATION NUMBER: US/09/383,586
;  CURRENT FILING DATE: 1999-08-26
;  NUMBER OF SEQ ID NOS: 38
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 31
;   LENGTH: 529
;   TYPE: PRT
;   ORGANISM: Mouse
US-09-383-586-31

  Query Match             90.9%;  Score 1560;  DB 3;  Length 529;
  Best Local Similarity   91.4%;  Pred. No. 1.4e-134;
  Matches  296;  Conservative    9;  Mismatches   15;  Indels    4;  Gaps    1;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    |||||  | | ||||||:||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              |||||||||||||||||||||||||:|| ||||||||||||||||||||:::|||
Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||
Db        117 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| ||||||||||:|||||||||||:||||||||||||||||
Db        177 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||:||||||||||||
Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||||| |
Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 3
US-09-823-038A-31
; Sequence 31, Application US/09823038A
; Patent No. 6797271
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT:  Sleeman, Matthew
;  APPLICANT:  Abernethy, Nevin
;  APPLICANT:  Onrust, Rene
;  APPLICANT:  Kumble, Anand
;  APPLICANT:  Murison, Greg
;  TITLE OF INVENTION: Compositions Isolated From Stromal Cells
;  TITLE OF INVENTION:  and Methods For Their Use
;  FILE REFERENCE: 11000.1037c3
;  CURRENT APPLICATION NUMBER: US/09/823,038A
;  CURRENT FILING DATE:  2001-07-09
;  NUMBER OF SEQ ID NOS: 61
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 31
;   LENGTH: 529
;   TYPE: PRT
;   ORGANISM: Mouse
US-09-823-038A-31

  Query Match             90.9%;  Score 1560;  DB 4;  Length 529;
  Best Local Similarity   91.4%;  Pred. No. 1.4e-134;
  Matches  296;  Conservative    9;  Mismatches   15;  Indels    4;  Gaps    1;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    |||||  | | ||||||:||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              |||||||||||||||||||||||||:|| ||||||||||||||||||||:::|||
Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||
Db        117 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| ||||||||||:|||||||||||:||||||||||||||||
Db        177 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 4
US-09-383-586-33
; Sequence 33, Application US/09383586
; Patent No. 6242419
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT: Sleeman, Matthew
;  APPLICANT: Abernethy, Nevin
;  APPLICANT: Onrust, Rene
;  APPLICANT: Kumble, Anand
;  APPLICANT: Murison, Greg
;  TITLE OF INVENTION: Compounds isolated from stromal cells
;  TITLE OF INVENTION: and methods for their use
;  FILE REFERENCE: 11000.1037c1
;  CURRENT APPLICATION NUMBER: US/09/383,586
;  CURRENT FILING DATE: 1999-08-26
;  NUMBER OF SEQ ID NOS: 38
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 33
;   LENGTH: 322
;   TYPE: PRT
;   ORGANISM: Human
US-09-383-586-33

  Query Match             84.9%;  Score 1458;  DB 3;  Length 322;
  Best Local Similarity   100.0%;  Pred. No. 1.5e-125;
  Matches  273;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy         52 PVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVN 111
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         50 PVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVN 109

Qy        112 YTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLK 171
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        110 YTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLK 169

Qy        172 CVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAIN 231
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Db        170 CVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAIN 229

Qy        232 ATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGR 291
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        230 ATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGR 289

Qy        292 HNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
                ||||||||||||||||||||||||||||||||
Db        290 HNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 322


RESULT 5
US-09-823-038A-33
; Sequence 33, Application US/09823038A
; Patent No. 6797271
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT:  Sleeman, Matthew
;  APPLICANT:  Abernethy, Nevin
;  APPLICANT:  Onrust, Rene
;  APPLICANT:  Kumble, Anand
;  APPLICANT:  Murison, Greg
;  TITLE OF INVENTION: Compositions Isolated From Stromal Cells
;  TITLE OF INVENTION:  and Methods For Their Use
;  FILE REFERENCE: 11000.1037c3
;  CURRENT APPLICATION NUMBER: US/09/823,038A
;  CURRENT FILING DATE:  2001-07-09
;  NUMBER OF SEQ ID NOS: 61
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 33
;   LENGTH: 322
;   TYPE: PRT
;   ORGANISM: Human
US-09-823-038A-33

  Query Match             84.9%;  Score 1458;  DB 4;  Length 322;
  Best Local Similarity   100.0%;  Pred. No. 1.5e-125;
  Matches  273;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy         52 PVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVN 111
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         50 PVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVN 109

Qy        112 YTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLK 171
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        110 YTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLK 169

Qy        172 CVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAIN 231
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        170 CVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAIN 229

Qy        232 ATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGR 291
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        230 ATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGR 289

Qy        292 HNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
                ||||||||||||||||||||||||||||||||
Db        290 HNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 322


RESULT 6
US-09-383-586-32
; Sequence 32, Application US/09383586
; Patent No. 6242419
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT: Sleeman, Matthew
;  APPLICANT: Abernethy, Nevin
;  APPLICANT: Onrust, Rene
;  APPLICANT: Kumble, Anand
;  APPLICANT: Murison, Greg
;  TITLE OF INVENTION: Compounds isolated from stromal cells
;  TITLE OF INVENTION: and methods for their use
;  FILE REFERENCE: 11000.1037c1
;  CURRENT APPLICATION NUMBER: US/09/383,586
;  CURRENT FILING DATE: 1999-08-26
;  NUMBER OF SEQ ID NOS: 38
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 32
;   LENGTH: 439
;   TYPE: PRT
;   ORGANISM: Mouse
US-09-383-586-32

  Query Match             60.3%;  Score 1035.5;  DB 3;  Length 439;
  Best Local Similarity   65.1%;  Pred. No. 1.3e-86;
  Matches  211;  Conservative    4;  Mismatches   14;  Indels   95;  Gaps    2;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
                || || |||    |||||  | | |||
Db          1 MTRSPALLL----LLLGALPSAEAAR-------------------------------- 22

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
                                                                    |||
Db         23 ----------------------------------------------------------DDI 25

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
                ||||||  ||  ||||||||||||||||||||||||||||||||||||||||||||||||||
Db         26 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 85

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
                || |||||| ||  ||:| |||||||||||:|||||||||||||:|||||||||||||
Db         86 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 145

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
                |||||||||||||||||||||||||||||||||||||||||||||:||||||||||||
Db        146 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 205

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
                |||||||||||||||||||| |
Db        206 QKFVVLPTGDVWSRPDGSYLNKLL 229


RESULT 7
US-09-823-038A-32
; Sequence 32, Application US/09823038A
; Patent No. 6797271
; GENERAL INFORMATION:
;  APPLICANT: Strachan, Lorna
;  APPLICANT:  Sleeman, Matthew
;  APPLICANT:  Abernethy, Nevin
;  APPLICANT:  Onrust, Rene
;  APPLICANT:  Kumble, Anand
;  APPLICANT:  Murison, Greg
;  TITLE OF INVENTION: Compositions Isolated From Stromal Cells
;  TITLE OF INVENTION:  and Methods For Their Use
;  FILE REFERENCE: 11000.1037c3
;  CURRENT APPLICATION NUMBER: US/09/823,038A
;  CURRENT FILING DATE:  2001-07-09
;  NUMBER OF SEQ ID NOS: 61
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 32
;   LENGTH: 439
;   TYPE: PRT
;   ORGANISM: Mouse
US-09-823-038A-32

  Query Match             60.3%;  Score 1035.5;  DB 4;  Length 439;
  Best Local Similarity   65.1%;  Pred. No. 1.3e-86;
  Matches  211;  Conservative    4;  Mismatches   14;  Indels   95;  Gaps    2;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
                || || |||    |||||  | | |||
Db          1 MTRSPALLL----LLLGALPSAEAAR-------------------------------- 22
```

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Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
                                                                    |||
Db         23 ----------------------------------------------------DDI 25

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||||
Db         26 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 85

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| ||||||||||||:||||||||||||:||||||||||||||||
Db         86 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 145

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||:||||||||||||||||||
Db        146 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 205

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||| |
Db        206 QKFVVLPTGDVWSRPDGSYLNKLL 229


RESULT 8
US-09-383-630-6
; Sequence 6, Application US/09383630A
; Patent No. 6265632
;   GENERAL INFORMATION:
;        APPLICANT: Avner Yayon et al.
;        TITLE OF INVENTION: ANIMAL MODEL FOR FIBROBLAST GROWTH
;                            FACTOR RECEPTOR ASSOCIATED
;                            CHONDRODYSPLASIA
;        NUMBER OF SEQUENCES: 18
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Mark M. Friedman c/o Anthony Castorina
;             STREET: 2001 Jefferson Davis Highway, Suite 207
;             CITY: Arlington
;             STATE: Virginia
;             COUNTRY: United States of America
;             ZIP: 22202
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
;             COMPUTER: Twinhead* Slimnote-890TX
;             OPERATING SYSTEM: MS DOS version 6.2,
;                               Windows version 3.11
;             SOFTWARE: Word for Windows version 2.0 converted
;                       to an ASCI file
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/383,630A
;             FILING DATE: 26-Aug-1999
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: <Unknown>
;             FILING DATE: <Unknown>
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Friedmam, Mark M.
;             REGISTRATION NUMBER: 33,883
;             REFERENCE/DOCKET NUMBER: 1402/2
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 972-3-5625553
;             TELEFAX: 972-3-5625554
;             TELEX: <Unknown>
;   INFORMATION FOR SEQ ID NO: 6:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 801
;             TYPE: amino acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-09-383-630-6

  Query Match             23.8%;  Score 408.5;  DB 3;  Length 801;
  Best Local Similarity   34.4%;  Pred. No. 9.2e-29;
  Matches  106;  Conservative   49;  Mismatches  114;  Indels   39;  Gaps   11;
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Qy         22 AAAARGPPKMADKVVPR----------QVA-RLGRTVRLQC-PVEGDPPPLTMWTKD 66
              | |   ||    :|| |             |||   | || | | |  | |   |:| ||
Db         16 AGATSEPPGPEQRVVRRAAEVPGPEPSQQEQVAFGSGDTVELSCHPPGGAPTGPTVWAKD 75

Qy         67 GRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGK 124
              | : :   |  | || |:|     ||||| |  : |     :  :::: | |  | |
Db         76 GTGLVAS-HRILVGPQRLQVLNASHEDAGVYSCQHRLTR---RVLCHFSVRVTDAPSSGD 131

Qy        125 ESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITW 184
              :  | | :    ||  :  |     |:| :| ::::| |  ::|| :| |:|:| | |:|
Db        132 DEDGEDVA----EDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISW 182

Qy        185 MKDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQR 241
              :|: :      |    : | ::|:|.:::: | | | ||| | |: |:|  || :||::|
Db        183 LKNGKEFRGEHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLER 242

Qy        242 TRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQ 301
              :  :|:|    | | |   |    | ||| || :| ||||| ||      : | :   |
Db        243 SPHRPILQAGLPANQTAILGSDVEFHCKVYSDAQPHIQWLKHVEV-----NGSKVGPDGT 297

Qy        302 KFV-VLPT 308
               :| || |
Db        298 PYVTVLKT 305


RESULT 9
US-09-383-630-3
; Sequence 3, Application US/09383630A
; Patent No. 6265632
;   GENERAL INFORMATION:
;        APPLICANT: Avner Yayon et al.
;        TITLE OF INVENTION: ANIMAL MODEL FOR FIBROBLAST GROWTH
;                            FACTOR RECEPTOR ASSOCIATED
;                            CHONDRODYSPLASIA
;        NUMBER OF SEQUENCES: 18
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Mark M. Friedman c/o Anthony Castorina
;             STREET: 2001 Jefferson Davis Highway, Suite 207
;             CITY: Arlington
;             STATE: Virginia
;             COUNTRY: United States of America
;             ZIP: 22202
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
;             COMPUTER: Twinhead* Slimnote-890TX
;             OPERATING SYSTEM: MS DOS version 6.2,
;                               Windows version 3.11
;             SOFTWARE: Word for Windows version 2.0 converted
;                       to an ASCI file
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/383,630A
;             FILING DATE: 26-Aug-1999
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: <Unknown>
;             FILING DATE: <Unknown>
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Friedmam, Mark M.
;             REGISTRATION NUMBER: 33,883
;             REFERENCE/DOCKET NUMBER: 1402/2
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 972-3-5625553
;             TELEFAX: 972-3-5625554
;             TELEX: <Unknown>
;   INFORMATION FOR SEQ ID NO: 3:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 806
;             TYPE: amino acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-383-630-3
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  Query Match             22.9%;  Score 393;  DB 3;  Length 806;
  Best Local Similarity   33.4%;  Pred. No. 2.5e-27;
  Matches  100;  Conservative   49;  Mismatches  116;  Indels   34;  Gaps   11;

Qy         22 AAAARGPPKMADKVVPRQVARL----GRTVRLQCPVEGDPP--PLTMWTKDGRTIHSGWS 75
              ||   ||        | |  :|    |  | | ||  |  |  | |:| |||  :
Db         35 AAEVPGPE-------PGQQEQLVFGSGDAVELSCPPPGGGPMGP-TVWVKDGTGLVPS-E 85

Qy         76 RFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGKESLGPDSSS 133
              |  | || |:|     ||:| | |  :  |     :  :::: | |  | :  | | :
Db         86 RVLVGPQRLQVLNASHEDSGAYSCRQRLTQ---RVLCHFSVRVTDAPSSGDDEDGEDEAE 142

Qy        134 GGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT- 192
                 |  :  |     |:| :| ::::| |  ::|| :| |:|:| | |:|:|: :
Db        143 DTGVDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISWLKNGREFRG 197

Qy        193 --RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTG 250
                |  : | ::|:| :::: | | | ||| | |: |:|  || :||::|:  :|:|
Db        198 EHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLERSPHRPILQA 257

Qy        251 THPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFV-VLPT 308
              | | |   |    | ||| || :| ||||| ||      : | :   | :| || |
Db        258 GLPANQTAVLGSDVEFHCKVYSDAQPHIQWLKHVEV-----NGSKVGPDGTPYVTVLKT 311


RESULT 10
US-09-173-151A-33
; Sequence 33, Application US/09173151A
; Patent No. 6326472
;  GENERAL INFORMATION:
;    APPLICANT:  Timans, Jacqueline C.
;    APPLICANT:  Debets, Johannes Eduard Maria
;    APPLICANT:  Antonius
;    APPLICANT:  Sana, Theodore R.
;    APPLICANT:  Bazan, J. Fernando
;    APPLICANT:  Kastelein, Robert A.
;    TITLE OF INVENTION:  Human Receptor Proteins; Related Reagents and Methods
;    NUMBER OF SEQUENCES:  36
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  DNAX Research Institute
;      STREET:  901 California Avenue
;      CITY:  Palo Alto
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94304-1104
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/173,151A
;      FILING DATE:  14-OCT-1998
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/065,776
;      FILING DATE:  17-NOV-1997
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/078,008
;      FILING DATE:  12-MAR-1998
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/081,883
;      FILING DATE:  15-APR-1998
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/095,987
;      FILING DATE:  10-AUG-1998
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/078,416
;      FILING DATE:  18-MAR-1998
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 60/062,066
;      FILING DATE:  15-OCT-1997
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Ching, Edwin P.
;      REGISTRATION NUMBER:  34,090
;      REFERENCE/DOCKET NUMBER:  DX0767X
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (650)852-9196
;      TELEFAX:  (650)496-1200
;  INFORMATION FOR SEQ ID NO:  33:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  802 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  not relevant
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  peptide
US-09-173-151A-33

  Query Match             22.3%;  Score 382.5;  DB 3;  Length 802;
  Best Local Similarity   31.3%;  Pred. No. 2.2e-26;
  Matches  109;  Conservative   50;  Mismatches  136;  Indels   53;  Gaps   12;

Qy          6 LLLLLLPPLLLGAFPPAAAARG------PPKMADKVVPRQ---VARLGRTVRLQCPVEGD 56
              ||| ||  ||     ||  :          | :|  :  ::      ||: ||| |   |
Db          3 LLLALLGVLLSVPGPPVLSLEASEEVELEPCLAPSLEQQEQELTVALGQPVRLCC---GR 59

Qy         57 PPPLTMWTKDGRTIHSGWSRFRVLPQG--------LKVKQVEREDAGVYVCKATNGFGSL 108
                  | |:|           |: | |       |::     |||| |:| |    ||:
Db         60 AERGGHWYKEGS---------RLAPAGRVRGWRGRLEIASFLPEDAGRYLCLAR---GSM 107

Qy        109 SV--NYTLVVLDDISPGKESLGPDSSSGGQEDPA---SQQWARPRFTQPSKMRRRVIARP 163
               |  | ||:  | ::   :   | |      ||:   |      | :| | :| ::: | |
Db        108 IVLQNLTLITGDSLTSSNDDEDPKS----HRDPSNRHSYPQQAPYWTHPQRMEKKLHAVP 163

Qy        164 VGSSVRLKCVASGHPRPDITWMKDDQAL---TRPEAAEPRKKKWTLSLKNLRPEDSGKYT 220
               |::|: :|  |:|:|  | | |:|| ||      |      | : |:| :::: | | | ||
Db        164 AGNTVKFRCPAAGNPTPTIRWLKDGQAFHGENRIGGIRLRHQHWSLVMESVVPSDRGTYT 223

Qy        221 CRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQW 280
              | | |  |:|    | :||::|:  :|:|    | |||   |      ||| || :| |||
Db        224 CLVENAVGSIRYNYLLDVLERSPHRPILQAGLPANTTAVVGSDVELLCKVYSDAQPHIQW 283

Qy        281 LKRV-----EYGAEG----RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
              || :      :|| |    :   | |:   :  ||   :| :   | |
Db        284 LKHIVINGSSFGAVGFPYVQVLKTADINSSEVEVLYLRNVSAEDAGEY 331


RESULT 11
US-07-640-029-1
; Sequence 1, Application US/07640029
; Patent No. 5229501
;  GENERAL INFORMATION:
;    APPLICANT:  Kiefer, Michael C.
;    APPLICANT:  Valenzuela, Pablo D.T.
;    APPLICANT:  Barr, Philip J.
;    TITLE OF INVENTION:  Expression and Use of Human Fibroblast
;    TITLE OF INVENTION:  Growth Factor Receptor
;    NUMBER OF SEQUENCES:  12
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Chiron Corporation
;      STREET:  4560 Horton Street
;      CITY:  Emeryville
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94608
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/640,029
;      FILING DATE:  19910111
;      CLASSIFICATION:  530
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  McClung, Barbara G.
```

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;      REGISTRATION NUMBER:  33,113
;      REFERENCE/DOCKET NUMBER:  CH-165
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  510-601-2708
;      TELEFAX:  510-655-3542
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  816 amino acids
;      TYPE:  AMINO ACID
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  peptide
US-07-640-029-1

  Query Match             22.1%;  Score 380;  DB 1;  Length 816;
  Best Local Similarity   30.0%;  Pred. No. 3.9e-26;
  Matches   89;  Conservative   53;  Mismatches  125;  Indels   30;  Gaps    8;

Qy          3 PSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTM 62
              |||| |           | |   | |   :     :   | ::|:| :  |   :
Db         23 PSPTL-----------PEQAQPWGAPVEVESF----LVHPGDLLQLRCRLRDDVQSIN- 65

Qy         63 WTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYYVCKATNGFGSLSVNYTLVVLDDISP 122
              | :||  :    :| |:  : ::|:     |:|:| |  ::  ||    |  | : |  |
Db         66 WLRDGVQLAES-NRTRITGEEVEVQDSVPADSGLYACVTSSPSGS-DTTYFSVNVSDALP 123

Qy        123 GKESLGPDSSSGGQE------DPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASG 176
                |    | |  :|      |        | :| | || ::: | |   :|: || :||
Db        124 SSEDDDDDDDSSSEEKKEKETDNTKPNPVAPYWTSPEKMEKKLHAVPAAKTVKFKCPSSG 183

Qy        177 HPRPDITWMKDDQALTRPE----AAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINA 232
                | | : |:|: :   :|:      : |   |:: : :: | | | ||| | |  |:||
Db        184 TPNPTLRWLKNGKEF-KPDHRIGGYKVRYATWSIIMDSVVPSDKGNYTCIVENEYGSINH 242

Qy        233 TYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAE 289
              ||:|:|::|:  :|:|    | | ||  |   | ||| || :| ||||| :|:|::
Db        243 TYQLDVVERSPHRPILQAGLPANKTVALGSNVEFMCKVYSDPQPHIQWLKHIEWGSK 299


RESULT 12
US-08-471-570-14
; Sequence 14, Application US/08471570
; Patent No. 5750371
;  GENERAL INFORMATION:
;    APPLICANT:  IGARASHI, Koichi
;    APPLICANT:  SENOO, Masaharu
;    APPLICANT:  WATANABE, Tatsuya
;    TITLE OF INVENTION:  PROTEIN, DNA AND USE THEREOF
;    NUMBER OF SEQUENCES:  18
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  DAVID G. CONLIN; DIKE, BRONSTEIN, ROBERTS &
;      ADDRESSEE:  CUSHMAN
;      STREET:  130 Water Street
;      CITY:  Boston
;      STATE:  Massachusetts
;      COUNTRY:  US
;      ZIP:  02109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/471,570
;      FILING DATE:  06-JUN-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/149,664
;      FILING DATE:
;      APPLICATION NUMBER:  US 07/743369
;      FILING DATE:  16-AUG-1991
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  LINEK, Ernest V
;      REGISTRATION NUMBER:  29822
;      REFERENCE/DOCKET NUMBER:  40897
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)523-3400
;      TELEFAX:  (617)523-6440
;      TELEX:  200291 STRE UR
;  INFORMATION FOR SEQ ID NO:  14:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  355 amino acids
;      TYPE:  amino acid
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-08-471-570-14

  Query Match             22.1%;  Score 379.5;  DB 1;  Length 355;
  Best Local Similarity   30.1%;  Pred. No. 1.3e-26;
  Matches   97;  Conservative   59;  Mismatches  129;  Indels   37;  Gaps   10;

Qy         20 PPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRV 79
              ||       |::        ||  | :: ::| ::      :  |||||  :| | :   |
Db         18 PPTKYQISQPEV-------YVAAPGESLEVRCLLK--DAAVISWTKDG--VHLGPNNRTV 66

Qy         80 L-PQGLKVKQVEREDAGVYYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED 138
              |  : |::|     |:|:| | |:    | :  : : | | || | :    :  : | ||
Db         67 LIGEYLQIKGATPRDSGLYACTASRTVDSETWYFMVNVTDAISSGDD----EDDTDGAED 122

Qy        139 PASQ---QWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT--- 192
                |:       | :|   || :|: | |  ::|: :| | |:| | : |:|: :
Db        123 FVSENSNNKRAPYWTNTEKMEKRLHAVPAANTVKFRCPAGGNPMPTMRWLKNGKEFKQEH 182

Qy        193 RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTH 252
              |    : | : |:| :::: | | | ||| | |  |:|| || :||::|:  :|:|
Db        183 RIGGYKVRNQHWSLIMESVVPSDKGNYTCVVENEYGSINHTYHLDVVERSPHRPILQAGL 242

Qy        253 PVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEG-------RHNSTIDVGG 300
              | | :   ||    | ||| || :| |||:| ||      || :|       :|:    :
Db        243 PANASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDGLPYLKVLKHSG---INS 299

Qy        301 QKFVVLPTGDVWSRPDGSYLNK 322
                    ||   :|     | |: |
Db        300 SNAEVLALFNVTEADAGEYICK 321


RESULT 13
US-08-471-570-6
; Sequence 6, Application US/08471570
; Patent No. 5750371
;  GENERAL INFORMATION:
;    APPLICANT:  IGARASHI, Koichi
;    APPLICANT:  SENOO, Masaharu
;    APPLICANT:  WATANABE, Tatsuya
;    TITLE OF INVENTION:  PROTEIN, DNA AND USE THEREOF
;    NUMBER OF SEQUENCES:  18
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  DAVID G. CONLIN; DIKE, BRONSTEIN, ROBERTS &
;      ADDRESSEE:  CUSHMAN
;      STREET:  130 Water Street
;      CITY:  Boston
;      STATE:  Massachusetts
;      COUNTRY:  US
;      ZIP:  02109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/471,570
;      FILING DATE:  06-JUN-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/149,664
;      FILING DATE:
```

; APPLICATION NUMBER: US 07/743369
; FILING DATE: 16-AUG-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: LINEK, Ernest V
; REGISTRATION NUMBER: 29822
; REFERENCE/DOCKET NUMBER: 40897
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617)523-3400
; TELEFAX: (617)523-6440
; TELEX: 200291 STRE UR
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 643 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-471-570-6

Query Match 22.1%; Score 379.5; DB 1; Length 643;
Best Local Similarity 30.1%; Pred. No. 3.1e-26;
Matches 97; Conservative 59; Mismatches 129; Indels 37; Gaps 10;

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Qy          20 PPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRV 79
               ||        |::         ||  | :: ::| ::      :  |||||  :| | :   |
Db          38 PPTKYQISQPEV-------YVAAPGESLEVRCLLK--DAAVISWTKDG--VHLGPNNRTV 86

Qy          80 L-PQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED 138
               |  : |::|     |:|:| | |:    | :  : : | | || | :    :  : | ||
Db          87 LIGEYLQIKGATPRDSGLYACTASRTVDSETWYFMVNVTDAISSGDD----EDDTDGAED 142

Qy         139 PASQ---QWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT--- 192
                 |:       | :|   || :|: | |  ::|: :| | |:| | : |:|: :
Db         143 FVSENSNNKRAPYWTNTEKMEKRLHAVPAANTVKFRCPAGGNPMPTMRWLKNGKEFKQEH 202

Qy         193 RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTH 252
               |    : | : |:| :::: | | | ||| | |  |:|| || :||::|:  :|:|
Db         203 RIGGYKVRNQHWSLIMESVVPSDKGNYTCVVENEYGSINHTYHLDVVERSPHRPILQAGL 262

Qy         253 PVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEG-------RHNSTIDVGG 300
               | | :   ||   | ||| || :| |||:| ||     || :|      :|:    :
Db         263 PANASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDGLPYLKVLKHSG---INS 319

Qy         301 QKFVVLPTGDVWSRPDGSYLNK 322
                     ||   :|      | |: |
Db         320 SNAEVLALFNVTEADAGEYICK 341
```

RESULT 14
US-08-471-570-8
; Sequence 8, Application US/08471570
; Patent No. 5750371
; GENERAL INFORMATION:
; APPLICANT: IGARASHI, Koichi
; APPLICANT: SENOO, Masaharu
; APPLICANT: WATANABE, Tatsuya
; TITLE OF INVENTION: PROTEIN, DNA AND USE THEREOF
; NUMBER OF SEQUENCES: 18
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: DAVID G. CONLIN; DIKE, BRONSTEIN, ROBERTS &
; ADDRESSEE: CUSHMAN
; STREET: 130 Water Street
; CITY: Boston
; STATE: Massachusetts
; COUNTRY: US
; ZIP: 02109
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/471,570
; FILING DATE: 06-JUN-1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/149,664
; FILING DATE:
; APPLICATION NUMBER: US 07/743369
; FILING DATE: 16-AUG-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: LINEK, Ernest V
; REGISTRATION NUMBER: 29822
; REFERENCE/DOCKET NUMBER: 40897
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617)523-3400
; TELEFAX: (617)523-6440
; TELEX: 200291 STRE UR
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 769 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-471-570-8

Query Match 22.1%; Score 379.5; DB 1; Length 769;
Best Local Similarity 30.1%; Pred. No. 4e-26;
Matches 97; Conservative 59; Mismatches 129; Indels 37; Gaps 10;

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Qy          20 PPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRV 79
               ||        |::         ||  | :: ::| ::      :  |||||  :| | :   |
Db          38 PPTKYQISQPEV-------YVAAPGESLEVRCLLK--DAAVISWTKDG--VHLGPNNRTV 86

Qy          80 L-PQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED 138
               |  : |::|     |:|:| | |:    | :  : : | | || | :    :  : | ||
Db          87 LIGEYLQIKGATPRDSGLYACTASRTVDSETWYFMVNVTDAISSGDD----EDDTDGAED 142

Qy         139 PASQ---QWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT--- 192
                 |:       | :|   || :|: | |  ::|: :| | |:| | : |:|: :
Db         143 FVSENSNNKRAPYWTNTEKMEKRLHAVPAANTVKFRCPAGGNPMPTMRWLKNGKEFKQEH 202

Qy         193 RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTH 252
               |    : | : |:| :::: | | | ||| | |  |:|| || :||::|:  :|:|
Db         203 RIGGYKVRNQHWSLIMESVVPSDKGNYTCVVENEYGSINHTYHLDVVERSPHRPILQAGL 262

Qy         253 PVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEG-------RHNSTIDVGG 300
               | | :   ||   | ||| || :| |||:| ||     || :|      :|:    :
Db         263 PANASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDGLPYLKVLKHSG---INS 319

Qy         301 QKFVVLPTGDVWSRPDGSYLNK 322
                     ||   :|      | |: |
Db         320 SNAEVLALFNVTEADAGEYICK 341
```

RESULT 15
US-08-451-822A-13
; Sequence 13, Application US/08451822A
; Patent No. 5863888
; GENERAL INFORMATION:
; APPLICANT: Dionne, Craig A
; APPLICANT: Crumley, Greg
; APPLICANT: Jaye, Michael C
; APPLICANT: Schlessinger, Joseph
; TITLE OF INVENTION: Fibroblast Growth Factor Receptors
; NUMBER OF SEQUENCES: 19
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Rhone-Poulenc Rorer Legal Department
; STREET: 500 Arcola Road
; CITY: Collegeville
; STATE: PA
; COUNTRY: USA
; ZIP: 19426
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS

```
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/451,822A
;      FILING DATE:  26-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 08/323,430
;      FILING DATE:  14-OCT-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/934,372
;      FILING DATE:  21-AUG-1992
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/549,587
;      FILING DATE:  06-JUL-1990
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Savitzky, Martin
;      REGISTRATION NUMBER:  29,699
;      REFERENCE/DOCKET NUMBER:  A0496E
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (610) 454-3816
;      TELEFAX:  (610) 454-3808
;  INFORMATION FOR SEQ ID NO:  13:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  821 amino acids
;      TYPE:  amino acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  peptide
US-08-451-822A-13

  Query Match             21.9%;  Score 375.5;  DB 2;  Length 821;
  Best Local Similarity   31.8%;  Pred. No. 1e-25;
  Matches   90;  Conservative   54;  Mismatches  112;  Indels   27;  Gaps    8;

Qy         20 PPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRV 79
              ||       |::         ||   | :: ::| ::     :  |||||  :| | :    |
Db         38 PPTKYQISQPEV-------YVAAPGESLEVRCLLK--DAAVISWTKDG--VHLGPNNRTV 86

Qy         80 L-PQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED 138
              |  : |::|     |:|:| | |:    | :  : : | | || | :    :  : | ||
Db         87 LIGEYLQIKGATPRDSGLYACTASRTVDSETWYFMVNVTDAISSGDD----EDDTDGAED 142

Qy        139 PASQ---QWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT--- 192
               |:        | :|   || :|: | |  ::|: :| | |:| | : |:|: :
Db        143 FVSENSNNKRAPYWTNTEKMEKRLHAVPAANTVKFRCPAGGNPMPTMRWLKNGKEFKQEH 202

Qy        193 RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTH 252
              |    : | : |:| :::: | | | ||| | |   |:|| || :||::|:  :|:|
Db        203 RIGGYKVRNQHWSLIMESVVPSDKGNYTCVVENEYGSINHTYHLDVVERSPHRPILQAGL 262

Qy        253 PVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEG 290
              | | :   ||   | ||| || :| |||:| ||     || :|
Db        263 PANASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDG 305


Search completed: February  2, 2005, 18:26:58
Job time : 29 secs
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GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 2, 2005, 18:17:51 ; Search time 18 Seconds
(without alignments)
1731.902 Million cell updates/sec

Title: US-10-613-413B-8
Perfect score: 1717
Sequence: 1 MTPSPLLLLLLPPLLLGAFP.....VLPTGDVWSRPDGSYLNKPL 324

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR_79:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	425.5	24.8	812	1	A36477	fibroblast growth
2	424.5	24.7	814	1	A39752	fibroblast growth
3	422	24.6	822	2	B49151	fibroblast growth
4	416	24.2	822	2	S19947	fibroblast growth
5	409	23.8	800	1	TVHU2F	fibroblast growth
6	408.5	23.8	800	2	A48991	heparin-binding gr
7	408.5	23.8	801	2	I55363	fibroblast growth
8	399.5	23.3	797	2	S38579	fibroblast growth
9	393.5	22.9	806	2	A35963	protein-tyrosine k
10	393	22.9	806	1	TVHUF3	fibroblast growth
11	389.5	22.7	819	1	TVCHFG	fibroblast growth
12	389.5	22.7	822	2	B54846	fibroblast growth
13	385.5	22.5	818	2	JC4058	fibroblast growth
14	383.5	22.3	824	2	S24108	protein-tyrosine k
15	382.5	22.3	802	1	TVHUF4	fibroblast growth
16	382	22.2	480	2	A56182	fibroblast growth
17	382	22.2	829	2	JC4583	fibroblast growth
18	381	22.2	480	2	B56182	fibroblast growth
19	380	22.1	816	2	A49151	fibroblast growth
20	379.5	22.1	769	2	S16236	fibroblast growth
21	379.5	22.1	821	1	TVMSBK	fibroblast growth
22	379.5	22.1	822	2	A45081	fibroblast growth
23	379.5	22.1	822	2	A41794	keratinocyte growt
24	379.5	22.1	823	2	B35963	protein-tyrosine k
25	375.5	21.9	821	1	TVHUF2	fibroblast growth
26	372.5	21.7	822	2	I49289	fibroblast growth
27	371.5	21.6	799	2	S18209	fibroblast growth
28	370.5	21.6	822	2	S29840	fibroblast growth
29	368	21.4	820	2	S17295	fibroblast growth
30	367.5	21.4	662	2	C40862	heparin-binding gr
31	367.5	21.4	822	1	TVHUFG	fibroblast growth
32	364.5	21.2	361	2	PN0020	fibroblast growth
33	363.5	21.2	832	2	JH0393	fibroblast growth
34	361.5	21.1	822	1	TVMSFG	fibroblast growth
35	360	21.0	824	2	S36439	fibroblast growth
36	352.5	20.5	813	1	A49123	fibroblast growth
37	324	18.9	353	2	S51242	heparin-binding fi
38	318	18.5	682	2	A35969	heparin-binding gr
39	315	18.3	713	2	I50128	fibroblast growth
40	307	17.9	707	2	A38429	keratinocyte growt
41	307	17.9	707	2	A54846	fibroblast growth
42	307	17.9	729	2	A56795	fibroblast growth
43	305	17.8	733	2	I49293	fibroblast growth
44	304	17.7	302	2	C36464	fibroblast growth
45	304	17.7	705	2	S51635	fibroblast growth

ALIGNMENTS

RESULT 1
A36477
fibroblast growth factor receptor A2 precursor - African clawed frog
N;Contains: fibroblast growth factor receptor A2, short splice form; protein-tyrosine kir
C;Species: Xenopus laevis (African clawed frog)
C;Date: 08-Mar-1991 #sequence_revision 19-Jan-1996 #text_change 09-Jul-2004
C;Accession: A36477; C39752
R;Musci, T.J.; Amaya, E.; Kirschner, M.W.
Proc. Natl. Acad. Sci. U.S.A. 87, 8365-8369, 1990
A;Title: Regulation of the fibroblast growth factor receptor in early Xenopus embryos.
A;Reference number: A36477; MUID:91045998; PMID:2172985
A;Accession: A36477
A;Molecule type: mRNA
A;Residues: 1-812 <MUS>
A;Cross-references: UNIPROT:P22182; GB:U24491; GB:M37201; NID:g857677; PIDN:AAA86868.1; I
A;Note: 45-Arg, 49-Thr, 50-Val, and 64-Ser also found
R;Friesel, R.; Dawid, I.B.
Mol. Cell. Biol. 11, 2481-2488, 1991
A;Title: cDNA cloning and developmental expression of fibroblast growth factor receptors
A;Reference number: A39752; MUID:91203867; PMID:1850097
A;Accession: C39752
A;Molecule type: mRNA
A;Residues: 1-30,119-189,'G',191-418,'L',420-636,'R',638-788,'V',790-812 <FRI>
A;Cross-references: GB:M62322; NID:g214899; PIDN:AAA49993.1; PID:g214900
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: alternative splicing; ATP; autophosphorylation; duplication; glycoprotein; gl
protein kinase
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-812/Product: fibroblast growth factor receptor A2, long splice form #status predicte
F;22-372/Domain: extracellular #status predicted <EXT>
F;22-30,119-812/Product: fibroblast growth factor receptor A2, short splice form #status
F;47-102/Domain: immunoglobulin homology <IM1>
F;125-132/Region: acidic
F;167-228/Domain: immunoglobulin homology <IM2>
F;266-339/Domain: immunoglobulin homology <IM3>
F;373-393/Domain: transmembrane #status predicted <TMM>
F;394-812/Domain: intracellular #status predicted <INT>
F;470-755/Domain: protein kinase homology <KIN>
F;478-486/Region: protein kinase ATP-binding motif
F;54-100,174-226,273-337/Disulfide bonds: #status predicted
F;76,116,133,177,223,236,260,292,313,326/Binding site: carbohydrate (Asn) (covalent) #sta
F;508,525,617/Active site: Lys, Glu, Asp #status predicted
F;622,635/Binding site: magnesium (Asn, Asp) #status predicted
F;648/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

Query Match 24.8%; Score 425.5; DB 1; Length 812;
Best Local Similarity 33.2%; Pred. No. 3.7e-23;
Matches 106; Conservative 61; Mismatches 123; Indels 29; Gaps 11;

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Qy          9 LLLPPLLLGAFPPAAAARGPPKMADKVVPR-------QVARLGRTVRLQCPVEGDPPPLT 61
              |||  :||||    : || |  : |:| |:         |: |  : ||| :  |   :
Db          7 LLLWGVLLGA--ALSVARPPSTLPDEVAPKTKTEVEPYSAQPGDRITLQCRLREDVQSIN 64
```

```
Qy          62 MWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTL---VVLD 118
               | |:|  : |  :| |:  : :::     || ||| |  |||    |  ||:   | :
Db          65 -WVKNGVQL-SETNRTRITGEEIQISNAGPEDNGVYAC-VTNG---PSRTYTVLCSVNVS 118

Qy         119 DISPGKESLGPDSSSGGQEDPASQQWA--RPRFTQPSKMRRRVIARPVGSSVRLKCVASG 176
               |  |  |    |  :   |: |::     || :: | || ::: | |   :|: :| |:|
Db         119 DALPSAEDDDEDDDNSSSEEKAAENSKPNRPLWSHPEKMEKKLHAVPAAKTVKFRCPANG 178

Qy         177 HPRPDITWMKDDQALTRPE---AAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINAT 233
               | | : |:|:::|  : :    : | : |:| : :: | | | ||| | |: ||||  |
Db         179 TPTPTLRWLKNNRAFQQDQRIGGYKVRSQTWSLIMDSVVPSDKGNYTCIVENKYGAINHT 238

Qy         234 YKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHN 293
               |::||::|:  :|:|    | ||:|  | |  | ||| || :| ||||: :|     :
Db         239 YQLDVVERSPHRPILQAGLPANTSVTVGTTAEFSCKVYSDPQPHIQWLRHIEI-----NG 293

Qy         294 STIDVGGQKFV-VLPTGDV 311
               | :   |  :| :| |  |
Db         294 SRVASDGFPYVEILKTAGV 312
```

RESULT 2
A39752
fibroblast growth factor receptor A1 precursor - African clawed frog
N;Contains: fibroblast growth factor receptor A1, short splice form; protein-tyrosine ki
C;Species: Xenopus laevis (African clawed frog)
C;Date: 07-Feb-1992 #sequence_revision 19-Jan-1996 #text_change 09-Jul-2004
C;Accession: A39752; B39752
R;Friesel, R.; Dawid, I.B.
Mol. Cell. Biol. 11, 2481-2488, 1991
A;Title: cDNA cloning and developmental expression of fibroblast growth factor receptors
A;Reference number: A39752; MUID:91203867; PMID:1850097
A;Accession: A39752
A;Molecule type: mRNA
A;Residues: 1-814 <FRI>
A;Cross-references: UNIPROT:Q91897; GB:M55163; NID:g214893; PIDN:AAA49990.1; PID:g214894
A;Accession: B39752
A;Molecule type: mRNA
A;Residues: 1-30,119-814 <FR2>
A;Cross-references: GB:M55163
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: alternative splicing; ATP; autophosphorylation; duplication; glycoprotein; g
protein kinase
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-814/Product: fibroblast growth factor receptor A1, long splice form #status experim
F;22-372/Domain: extracellular #status predicted <EXT>
F;22-30,119-814/Product: fibroblast growth factor receptor A1, short splice form #status
F;47-102/Domain: immunoglobulin homology <IM1>
F;125-132/Region: acidic
F;167-228/Domain: immunoglobulin homology <IM2>
F;266-339/Domain: immunoglobulin homology <IM3>
F;373-393/Domain: transmembrane #status predicted <TMM>
F;394-814/Domain: intracellular #status predicted <INT>
F;472-757/Domain: protein kinase homology <KIN>
F;480-488/Region: protein kinase ATP-binding motif
F;54-100,174-226,273-337/Disulfide bonds: #status predicted
F;76,116,133,177,223,236,260,292,313,326/Binding site: carbohydrate (Asn) (covalent) #st
F;510,527,619/Active site: Lys, Glu, Asp #status predicted
F;624,637/Binding site: magnesium (Asn, Asp) #status predicted
F;650/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicte

  Query Match 24.7%; Score 424.5; DB 1; Length 814;
  Best Local Similarity 32.5%; Pred. No. 4.4e-23;
  Matches 103; Conservative 62; Mismatches 127; Indels 25; Gaps 10;

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Qy           9 LLLPPLLLGAFPPAAAARGPPKMADKVVPR-------QVARLGRTVRLQCPVEGDPPPLT 61
               |||  :||||    : || |  : |:| |:          || | || ||| :  |   ::
Db           7 LLLWGVLLGA--ALSVARPPSTLPDEVAPKTKTEVEPYSARPGDTVTLQCRLREDVQSIS 64

Qy          62 MWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDIS 121
               | |:|  :    :| |:  : :::     || |:| |      |: :| ::: |  |
Db          65 -WVKNGVQLLE-TNRTRITGEEIQISNAGPEDNGLYACVTIGPSGTYTVLFSINV-SDAQ 121

Qy         122 PGKESLGPDSSSGGQEDPASQQWA--RPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPR 179
               |  |    |  :   |: ||:     || :: | || ::: | |   :|: :| |:| |
Db         122 PSAEDDDDDDNSSSEEKASENSKPNRPFWSHPEKMEKKLHAVPAAKTVKFRCPANGTPS 181

Qy         180 PDITWMKDDQALTRPE----AAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYK 235
               | : |:|: :   ||:      : | : |:| : :: | | | ||| | |: | :| ||:
Db         182 PALRWLKNGKEF-RPDQRIGGYKVRSQTWSLIMDSVVPSDKGNYTCIVENKYGTLNHTYQ 240

Qy         236 VDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNST 295
               :||::|:  :|:|    | ||:|  | |  | ||| || :| ||||: :|      : |
Db         241 LDVVERSPHRPILQAGLPANTSVTVGSTAEFSCKVYSDPQPHIQWLRHIEI-----NGSR 295

Qy         296 IDVGGQKFV-VLPTGDV 311
               :   |  :| :| |  |
Db         296 VASDGFPYVEILKTAGV 312
```

RESULT 3
B49151
fibroblast growth factor receptor 4 - Iberian ribbed newt
C;Species: Pleurodeles waltlii (Iberian ribbed newt)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: B49151
R;Shi, D.L.; Feige, J.J.; Riou, J.F.; DeSimone, D.W.; Boucaut, J.C.
Development 116, 261-273, 1992
A;Title: Differential expression and regulation of two distinct fibroblast growth factor
A;Reference number: A49151; MUID:93130775; PMID:1483392
A;Accession: B49151
A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-822 <SHI>
A;Cross-references: UNIPROT:Q91288; GB:X65059; NID:g64252; PIDN:CAA46192.1; PID:g64253
A;Experimental source: tail-bud
A;Note: sequence extracted from NCBI backbone (NCBIN:122598, NCBIP:122599)
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; growth factor receptor
F;283-354/Domain: immunoglobulin homology <IMM>
F;484-769/Domain: protein kinase homology <KIN>
F;492-500/Region: protein kinase ATP-binding motif

  Query Match 24.6%; Score 422; DB 2; Length 822;
  Best Local Similarity 32.0%; Pred. No. 6.8e-23;
  Matches 110; Conservative 66; Mismatches 134; Indels 34; Gaps 12;

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Qy           2 TPSPLLLLLLPPLLLGAFPPAAAARGPPK-----MADKVVPRQVARL---GRTVRLQCPV 53
               |  || |||    |   ||   : ||  |:     :|: |   :   |   |  :|| |
Db          15 TTRPLALLLCGLL---AFSALSCARTLPEGRKANLAELVSEEEEHFLLDPGNALRLFC-- 69

Qy          54 EGDPPPLTMW-TKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNY 112
               : :   :  | |:  |  | |  | |:    |::  |  ||:|:|:|    | | :  |:
Db          70 DTNQTTIVNWYTESTRLQHGG--RIRLTDTVLEIADVTYEDSGLYLC-VVPGTGHILRNF 126

Qy         113 TLVVLDDISPG---KESLGPDSSSG--GQEDPASQQWARPRFTQPSKMRRRVIARPVGSS 167
               |: |:|  :: |    |  | : |:|  |:: | |  :  | ::|| :| ::: | | |::
Db         127 TISVVDSLASGDDDDEDHGREDSAGDMGEDPPYSTSYRAPFWSQPQRMDKKLYAVPAGNT 186

Qy         168 VRLKCVASGHPRPDITWMKDDQAL---TRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVS 224
               |: :| ::|:| | | |:|: :      |      | : |:| :::: | | | ||| |
Db         187 VKFRCPSAGNPTPGIRWLKNGREFGGEHRIGGIRLRHQHWSLVMESVVPSDRGNYTCLVE 246

Qy         225 NRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRV 284
               |: |:|: :| :||::|:  :|:|    | |||    |    | ||| || :| ||||| :
Db         247 NKFGSISYSYLLDVLERSPHRPILQAGLPANTTAMLGSDVQFFCKVYSDAQPHIQWLKHI 306

Qy         285 E-----YGAEG----RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
               |     || :|    :   | |:   :  ||   :|     | |
Db         307 EVNGSRYGPDGVPFVQVLKTADINSSEVEVLYLHNVSFEDAGEY 350
```

RESULT 4
S19947
fibroblast growth factor receptor - Iberian ribbed newt

```
C;Species: Pleurodeles waltlii (Iberian ribbed newt)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: S19947
R;Shi, D.L.; Feige, J.J.; Riou, J.F.; DeSimone, D.W.; Boucaut, J.C.
submitted to the EMBL Data Library, March 1992
A;Description: Receptores during early development of the urodele Pleurodeles waltlii.
A;Reference number: S19947
A;Accession: S19947
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-822 <SHI>
A;Cross-references: UNIPROT:Q91288; EMBL:X65059
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; growth factor receptor
F;283-354/Domain: immunoglobulin homology <IMM>
F;484-769/Domain: protein kinase homology <KIN>
F;492-500/Region: protein kinase ATP-binding motif

  Query Match             24.2%;  Score 416;  DB 2;  Length 822;
  Best Local Similarity   31.7%;  Pred. No. 1.9e-22;
  Matches  109;  Conservative   66;  Mismatches  135;  Indels   34;  Gaps   12;

Qy          2 TPSPLLLLLLPPLLLGAFPPAAAARGPPK-----MADKVVPRQVARL---GRTVRLQCPV 53
              |  || |||   |   ||   : ||  |:     :| | :  |   |  :|| |
Db         15 TTRPLALLLCGLL---AFSALSCARTLPEGRKANLAELVSEEEEHFLLDPGNALRLFC-- 69

Qy         54 EGDPPPLTMW-TKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNY 112
              : :   : | |:  |  | |  | |:     ::  |  ||:|:|:|    | | :  |:
Db         70 DTNQTTIVNWYTESTRLQHGG--RIRLTDTVQEIADVTYEDSGLYLC-VVPGTGHILRNF 126

Qy        113 TLVVLDDISPG---KESLGPDSSSG--GQEDPASQQWARPRFTQPSKMRRRVIARPVGSS 167
              |: |:|  :: |    |  | : |:|  |:: | |  :  | ::|| :| ::: | | |::
Db        127 TISVVDSLASGDDDDEDHGREDSAGDMGEDPPYSTSYRAPFWSQPQRMDKKLYAVPAGNT 186

Qy        168 VRLKCVASGHPRPDITWMKDDQAL---TRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVS 224
              |: :| ::|:| | | |:|: :      |      | : |:| :::: | | | ||| |
Db        187 VKFRCPSAGNPTPGIRWLKNGREFGGEHRIGGIRLRHQHWSLVMESVVPSDRGNYTCLVE 246

Qy        225 NRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRV 284
              |: |:|: :| :||::|:  :|:|    | |||   |    | ||| || :| ||||| :
Db        247 NKFGSISYSYLLDVLERSPHRPILQAGLPANTTAMLGSDVQFFCKVYSDAQPHIQWLKHI 306

Qy        285 E-----YGAEG----RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
              |     || :|    :   | |:   :  ||   :|      | |
Db        307 EVNGSRYGPDGVPFVQVLKTADINSSEVEVLYLHNVSFEDAGEY 350


RESULT 5
TVHU2F
fibroblast growth factor receptor flg-2 precursor - human
N;Contains: protein-tyrosine kinase (EC 2.7.1.112) flg-2
C;Species: Homo sapiens (man)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 16-Jun-2000
C;Accession: A60350; S21843
R;Avivi, A.; Zimmer, Y.; Yayon, A.; Yarden, Y.; Givol, D.
Oncogene 6, 1089-1092, 1991
A;Title: Flg-2, a new member of the family of fibroblast growth factor receptors.
A;Reference number: A60350; MUID:91296390; PMID:1648703
A;Accession: A60350
A;Molecule type: mRNA
A;Residues: 1-800 <AVI>
A;Cross-references: EMBL:X58255; NID:g31382; PIDN:CAA41209.1; PID:g31383
A;Experimental source: keratinocytes
C;Comment: This may be a receptor for keratinocyte growth factor.
C;Genetics:
A;Gene: GDB:FGFR2; JWS; CFD1; KGF; FLG2
A;Cross-references: GDB:127273; OMIM:176943
A;Map position: 10q25.3-10q26
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor receptor;
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-800/Product: fibroblast growth factor receptor flg-2 #status predicted <MAT>
F;22-369/Domain: extracellular #status predicted <EXT>
F;131-137/Region: acidic
F;262-335/Domain: immunoglobulin homology <IMM>
F;370-390/Domain: transmembrane #status predicted <TMM>
F;391-800/Domain: intracellular #status predicted <INT>
F;464-749/Domain: protein kinase homology <KIN>
F;472-480/Region: protein kinase ATP-binding motif
F;59-107,170-222,269-333/Disulfide bonds: #status predicted
F;96,219,256,288,309,322/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;502,519,611/Active site: Lys, Glu, Asp #status predicted
F;616,629/Binding site: magnesium (Asn, Asp) #status predicted
F;642/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

  Query Match             23.8%;  Score 409;  DB 1;  Length 800;
  Best Local Similarity   34.3%;  Pred. No. 5.8e-22;
  Matches  103;  Conservative   48;  Mismatches  111;  Indels   38;  Gaps   10;

Qy         22 AAAARGPPKMADKVVPR-------------QVA-RLGRTVRLQC-PVEGDPPPLTMWTKD 66
              | |   ||    :|| |               |||   | || | | |  | |   |:| ||
Db         16 AGATSEPPGPEQRVVRRAAEVPGPEPSQQEQVAFGSGDTVELSCHPPGGAPTGPTVWAKD 75

Qy         67 GRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGK 124
              |  : :   |  | || |:|     ||||| |  : |     :  :::: | |  | |
Db         76 GTGLVAS-HRILVGPQRLQVLNASHEDAGVYSCQHRLTR---RVLCHFSVRVTDAPSSGD 131

Qy        125 ESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITW 184
              :  | | :    ||  :  |      |:| :| :::: | |  ::|| :| |:|:| | |:|
Db        132 DEDGEDVA----EDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISW 182

Qy        185 MKDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQR 241
              :|: :      |    : | ::|:| :::: | | | ||| | |: |:|  || :||::|
Db        183 LKNGKEFRGQHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLER 242

Qy        242 TRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEGRHNSTI 296
              :  :|:|    | | |   |    | ||| || :| ||||| ||      | :|    |:
Db        243 SPHRPILQAGLPANQTAILGSDVEFHCKVYSDAQPHIQWLKHVEVNGSKVGPDGTPYDTV 302


RESULT 6
A48991
heparin-binding growth factor receptor - mouse
N;Alternate names: HBGF receptor
N;Contains: protein-tyrosine kinase (EC 2.7.1.112)
C;Species: Mus musculus (house mouse)
C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A48991
R;Katoh, O.; Hattori, Y.; Sasaki, H.; Sakamoto, H.; Fujimoto, K.; Fujii, T.; Sugimura, T.
Cancer Res. 53, 1136-1141, 1993
A;Title: Isolation of the complementary DNA encoding a mouse heparin-binding growth factc
A;Reference number: A48991; MUID:93177694; PMID:8382556
A;Accession: A48991
A;Status: preliminary
A;Molecule type: nucleic acid
A;Residues: 1-800 <KAT>
A;Cross-references: UNIPROT:Q61851; GB:S56291; NID:g298329; PIDN:AAB25535.1; PID:g298330
A;Experimental source: brain
A;Note: sequence extracted from NCBI backbone (NCBIN:126536, NCBIP:126537)
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; growth factor receptor; heparin binding; phosphotransferase; tyrosine-sp
F;262-335/Domain: immunoglobulin homology <IMM>
F;464-749/Domain: protein kinase homology <KIN>
F;472-480/Region: protein kinase ATP-binding motif

  Query Match             23.8%;  Score 408.5;  DB 2;  Length 800;
  Best Local Similarity   34.4%;  Pred. No. 6.3e-22;
  Matches  106;  Conservative   49;  Mismatches  114;  Indels   39;  Gaps   11;

Qy         22 AAAARGPPKMADKVVPR-------------QVA-RLGRTVRLQC-PVEGDPPPLTMWTKD 66
              | |   ||    :|| |               |||   | || | | |  | |   |:| ||
Db         16 AGATSEPPGPEQRVVRRAAEVPGPEPSQQEQVAFGSGDTVELSCHPPGGAPTGPTVWAKD 75

Qy         67 GRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGK 124
              |  : :   |  | || |:|     ||||| |  : |     :  :::: | |  | |
Db         76 GTGLVAS-HRILVGPQRLQVLNASHEDAGVYSCQHRLTR---RVLCHFSVRVTDAPSSGD 131
```

```
Qy        125 ESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITW 184
Db        132 DEDGEDVA----EDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISW 182

Qy        185 MKDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQR 241
Db        183 LKNGKEFRGEHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLER 242

Qy        242 TRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQ 301
Db        243 SPHRPILQAGLPANQTAILGSDVEFHCKVYSDAQPHIQWLKHVEV-----NGSKVGPDGT 297

Qy        302 KFV-VLPT 308
Db        298 PYVTVLKT 305
```

RESULT 7
I55363
fibroblast growth factor receptor 3 - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 09-Jul-2004
C;Accession: I55363; B53627
R;Ornitz, D.M.; Leder, P.
J. Biol. Chem. 267, 16305-16311, 1992
A;Title: Ligand specificity and heparin dependence of fibroblast growth factor receptors
A;Reference number: I55363; MUID:92355591; PMID:1379594
A;Accession: I55363
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-801 <RES>
A;Cross-references: UNIPROT:Q61851; GB:M81342; NID:g199144; PIDN:AAA39535.1; PID:g199145
R;Chellaiah, A.T.; McEwen, D.G.; Werner, S.; Xu, J.; Ornitz, D.M.
J. Biol. Chem. 269, 11620-11627, 1994
A;Title: Fibroblast growth factor receptor (FGFR) 3. Alternative splicing in immunoglobu
A;Reference number: A53627; MUID:94209351; PMID:7512569
A;Accession: B53627
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 242-364 <CHE>
A;Cross-references: GB:L26492
C;Genetics:
A;Gene: mFR3
A;Introns: 304/3; 353/1
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; growth factor receptor
F;262-335/Domain: immunoglobulin homology <IMM>
F;464-750/Domain: protein kinase homology <KIN>
F;472-480/Region: protein kinase ATP-binding motif

Query Match 23.8%; Score 408.5; DB 2; Length 801;
Best Local Similarity 34.4%; Pred. No. 6.3e-22;
Matches 106; Conservative 49; Mismatches 114; Indels 39; Gaps 11;

```
Qy         22 AAAARGPPKMADKVVPR-------------QVA-RLGRTVRLQC-PVEGDPPPLTMWTKD 66
Db         16 AGATSEPPGPEQRVVRRAAEVPGPEPSQQEQVAFGSGDTVELSCHPPGGAPTGPTVWAKD 75

Qy         67 GRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGK 124
Db         76 GTGLVAS-HRILVGPQRLQVLNASHEDAGVYSCQHRLTR---RVLCHFSVRVTDAPSSGD 131

Qy        125 ESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITW 184
Db        132 DEDGEDVA----EDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISW 182

Qy        185 MKDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQR 241
Db        183 LKNGKEFRGEHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLER 242

Qy        242 TRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQ 301
Db        243 SPHRPILQAGLPANQTAILGSDVEFHCKVYSDAQPHIQWLKHVEV-----NGSKVGPDGT 297

Qy        302 KFV-VLPT 308
Db        298 PYVTVLKT 305
```

RESULT 8
S38579
fibroblast growth factor receptor 3 - Iberian ribbed newt (fragment)
C;Species: Pleurodeles waltlii (Iberian ribbed newt)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 09-Jul-2004
C;Accession: S38579
R;Shi, D.L.; Fromentoux, V.; Launay, C.; Umbhauer, M.; Boucaut, J.C.
submitted to the EMBL Data Library, November 1993
A;Description: Expression of FGFR-3 in amphibian embryos.
A;Reference number: S38579
A;Accession: S38579
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-797 <SHI>
A;Cross-references: UNIPROT:Q91287; EMBL:X75603
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; growth factor receptor
F;258-331/Domain: immunoglobulin homology <IMM>
F;456-741/Domain: protein kinase homology <KIN>
F;464-472/Region: protein kinase ATP-binding motif

Query Match 23.3%; Score 399.5; DB 2; Length 797;
Best Local Similarity 33.5%; Pred. No. 2.8e-21;
Matches 95; Conservative 43; Mismatches 97; Indels 49; Gaps 8;

```
Qy         44 GRTVRLQCPVEGDPPPLTMWTKDGRTI------HSGWSRFRVLPQGLKVKQVEREDAGVY 97
Db         50 GDTIELSCTTPGSSVSV-VWFKDGISVDPPTWSHTG-------QKLLKIINVSYDDSGVY 101

Qy         98 VCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRR 157
Db        102 SCKARQSSEVLR-NVTVRVTD---------SPSSGDDEDDDEESESANAPKFTRPEWMEK 151

Qy        158 RVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT---RPEAAEPRKKKWTLSLKNLRPE 214
Db        152 KLLAVPAANTVRFRCPAAGKPTPSITWLKNGKEFKGEHRIGGIKLRHQQWSLVMESVVPS 211

Qy        215 DSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDV 274
Db        212 DRGNYTCVVANKYGTIRETYTLDVLERTPHRPILQAGFRSNKTVVVGSDVEFHCKVYSDA 271

Qy        275 KPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGS 318
Db        272 QPHIQWLKHVE------------VNGSKF----------GPDGN 293
```

RESULT 9
A35963
protein-tyrosine kinase (EC 2.7.1.112) cek2 precursor - chicken
C;Species: Gallus gallus (chicken)
C;Date: 09-Nov-1990 #sequence_revision 09-Nov-1990 #text_change 09-Jul-2004
C;Accession: A35963
R;Pasquale, E.B.
Proc. Natl. Acad. Sci. U.S.A. 87, 5812-5816, 1990
A;Title: A distinctive family of embryonic protein-tyrosine kinase receptors.
A;Reference number: A35963; MUID:90332672; PMID:2165604
A;Accession: A35963
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-806 <PAS>
A;Cross-references: UNIPROT:P18460; GB:M35195; NID:g211442; PIDN:AAA48664.1; PID:g211443
C;Genetics:
A;Gene: cek2
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor receptor;
F;1-23/Domain: signal sequence #status predicted <SIG>

```
F;24-806/Product: protein-tyrosine kinase cek2 #status predicted <MAT>
F;24-368/Domain: extracellular #status predicted <EXT>
F;131-138/Region: acidic
F;262-335/Domain: immunoglobulin homology <IMM>
F;369-389/Domain: transmembrane #status predicted <TMM>
F;390-806/Domain: intracellular #status predicted <INT>
F;464-749/Domain: protein kinase homology <KIN>
F;472-480/Region: protein kinase ATP-binding motif
F;61-107,170-222,269-333/Disulfide bonds: #status predicted
F;96,219,256,288,309,322/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;502,519,611/Active site: Lys, Glu, Asp #status predicted
F;616,629/Binding site: magnesium (Asn, Asp) #status predicted
F;642/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicte

  Query Match             22.9%;  Score 393.5;  DB 2;  Length 806;
  Best Local Similarity   32.8%;  Pred. No. 7.7e-21;
  Matches   98;  Conservative   52;  Mismatches  118;  Indels   31;  Gaps    9;

Qy         16 LGAFPPAAAARGPPK---MADKVVPRQVARL-------GRTVRLQCPVEGDPPPLTMWTK 65
              :||  |||  ||  :    | : :  : | |       | |: | |  :     :  | |
Db         17 VGAL-PAARRRGAERSGGQAAEYLRSETAFLEELVFGSGDTIELSCNTQSSSVSV-FWFK 74

Qy         66 DGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKE 125
              ||  |    :|  :  : ||:  |  :|:|:| ||  :    |  |:|: | |
Db         75 DGIGIAPS-NRTHIGQKLLKIINVSYDDSGLYSCKPRHSNEVLG-NFTVRVTD------- 125

Qy        126 SLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWM 185
               | |     :|  |:    | :|:| || ::::| |  ::|| :| | |:| | | |:
Db        126 --SPSSGDDEDDDDESEDTGVPFWTRPDKMEKKLLAVPAANTVRFRCPAGGNPTPTIYWL 183

Qy        186 KDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRT 242
              |: :      |    : | ::|:| :::: | | | ||| | |: | |  ||::||::|:
Db        184 KNGKEFKGEHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKYGNIRHTYQLDVLERS 243

Qy        243 RSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEGRHNSTI 296
                :|:|    | | ||  |    | ||| || :| |||| ||      || :|    |:
Db        244 PHRPILQAGLPANQTVVVGSNVEFHCKVYSDAQPHIQWLKHVEVNGSKYGPDGTPYVTV 302


RESULT 10
TVHUF3
fibroblast growth factor receptor 3 precursor - human
N;Contains: protein-tyrosine kinase (EC 2.7.1.112)
C;Species: Homo sapiens (man)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: A38576; A55273; E38269; I51880
R;Keegan, K.; Johnson, D.E.; Williams, L.T.; Hayman, M.J.
Proc. Natl. Acad. Sci. U.S.A. 88, 1095-1099, 1991
A;Title: Isolation of an additional member of the fibroblast growth factor receptor fami
A;Reference number: A38576; MUID:91142118; PMID:1847508
A;Accession: A38576
A;Molecule type: mRNA
A;Residues: 1-806 <KEE>
A;Cross-references: UNIPROT:P22607; GB:M58051; NID:g182568; PIDN:AAA52450.1; PID:g182569
R;Thompson, L.M.; Plummer, S.; Schalling, M.; Altherr, M.R.; Gusella, J.F.; Housman, D.E
Genomics 11, 1133-1142, 1991
A;Title: A gene encoding a fibroblast growth factor receptor isolated from the Huntingto
A;Reference number: A55273; MUID:92147110; PMID:1664411
A;Accession: A55273
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 76-394,'V',396-806 <THO>
A;Cross-references: GB:M64347; NID:g182564; PIDN:AAA58470.1; PID:g182565
A;Note: sequence extracted from NCBI backbone (NCBIP:80296)
R;Partanen, J.; Maekelae, T.P.; Alitalo, R.; Lehvaeslaiho, H.; Alitalo, K.
Proc. Natl. Acad. Sci. U.S.A. 87, 8913-8917, 1990
A;Title: Putative tyrosine kinases expressed in K-562 human leukemia cells.
A;Reference number: A38268; MUID:91062389; PMID:2247464
A;Accession: E38269
A;Molecule type: mRNA
A;Residues: 619-675 <PAR>
A;Cross-references: GB:M37782
R;Bellus, G.A.; Hefferon, T.W.; Ortiz de Luna, R.I.; Hecht, J.T.; Horton, W.A.; Machado,
Am. J. Hum. Genet. 56, 368-373, 1995
A;Title: Achondroplasia is defined by recurrent G380R mutations of FGFR3.
A;Reference number: I51880; MUID:95150025; PMID:7847369
A;Accession: I51880
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 361-379,'R',381-415 <RES>
A;Cross-references: GB:S76733; NID:g914201; PIDN:AAB33323.1; PID:g914202
A;Note: this sequence represents a mutant form associated with achondroplasia
C;Genetics:
A;Gene: GDB:FGFR3
A;Cross-references: GDB:127526; OMIM:100800; OMIM:134934
A;Map position: 4p16.3-4p16.3
C;Function:
A;Description: receptor for both acidic and basic fibroblast growth factors
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor receptor;
F;1-22/Domain: signal sequence #status predicted <SIG>
F;23-806/Product: fibroblast growth factor receptor 3 #status predicted <MAT>
F;23-375/Domain: extracellular #status predicted <EXT>
F;133-139/Region: acidic
F;268-341/Domain: immunoglobulin homology <IMM>
F;376-396/Domain: transmembrane #status predicted <TMM>
F;397-806/Domain: intracellular #status predicted <INT>
F;470-755/Domain: protein kinase homology <KIN>
F;478-486/Region: protein kinase ATP-binding motif
F;61-109,176-228,275-339/Disulfide bonds: #status predicted
F;98,225,262,294,315,328/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;508,525,617/Active site: Lys, Glu, Asp #status predicted
F;622,635/Binding site: magnesium (Asn, Asp) #status predicted
F;648/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

  Query Match             22.9%;  Score 393;  DB 1;  Length 806;
  Best Local Similarity   33.4%;  Pred. No. 8.4e-21;
  Matches  100;  Conservative   49;  Mismatches  116;  Indels   34;  Gaps   11;

Qy         22 AAAARGPPKMADKVVPRQVARL----GRTVRLQCPVEGDPP--PLTMWTKDGRTIHSGWS 75
              ||   ||         | |  :|    |  | | ||  |  |  | |:| |||  :
Db         35 AAEVPGPE-------PGQQEQLVFGSGDAVELSCPPPGGGPMGP-TVWVKDGTGLVPS-E 85

Qy         76 RFRVLPQGLKVKQVEREDAGVYVC--KATNGFGSLSVNYTLVVLDDISPGKESLGPDSSS 133
              |  | || |:|     ||:| | |  : |     :  :::: | |  | | :  | | :
Db         86 RVLVGPQRLQVLNASHEDSGAYSCRQRLTQ---RVLCHFSVRVTDAPSSGDDEDGEDEAE 142

Qy        134 GGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT- 192
                 |  :  |      |:| :| ::::| |   ::|| :| |:|:| | |:|:|: :
Db        143 DTGVDTGAPYW-----TRPERMDKKLLAVPAANTVRFRCPAAGNPTPSISWLKNGREFRG 197

Qy        193 --RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTG 250
                 |    : | ::|:| :::: | | | ||| | |: |:|  || :||::|:  :|:|
Db        198 EHRIGGIKLRHQQWSLVMESVVPSDRGNYTCVVENKFGSIRQTYTLDVLERSPHRPILQA 257

Qy        251 THPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFV-VLPT 308
              | | |   |    | ||| || :| ||||| ||      : | :   |  :| || |
Db        258 GLPANQTAVLGSDVEFHCKVYSDAQPHIQWLKHVEV-----NGSKVGPDGTPYVTVLKT 311


RESULT 11
TVCHFG
fibroblast growth factor receptor 1 precursor - chicken
N;Alternate names: basic fibroblast growth factor receptor
N;Contains: protein-tyrosine kinase (EC 2.7.1.112) cek1
C;Species: Gallus gallus (chicken)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: A41345; A33908
R;Lee, P.L.; Johnson, D.E.; Cousens, L.S.; Fried, V.A.; Williams, L.T.
Science 245, 57-60, 1989
A;Title: Purification and complementary DNA cloning of a receptor for basic fibroblast gr
A;Reference number: A41345; MUID:89298406; PMID:2544996
A;Accession: A41345
A;Status: nucleic acid sequence not shown; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-819 <LEE>
```

```
A;Cross-references: UNIPROT:P21804
A;Note: part of the sequence was confirmed by protein sequencing
R;Pasquale, E.B.; Singer, S.J.
Proc. Natl. Acad. Sci. U.S.A. 86, 5449-5453, 1989
A;Title: Identification of a developmentally regulated protein-tyrosine kinase by using
A;Reference number: A33908; MUID:89315814; PMID:2473471
A;Accession: A33908
A;Molecule type: mRNA
A;Residues: 1-89,'A',91-685,'M',687-819 <PAS>
A;Cross-references: GB:M24637
A;Note: this protein is expressed in embryonic tissues and, at low levels, in adult brai
C;Genetics:
A;Gene: cek1
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor receptor;
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-819/Product: fibroblast growth factor receptor 1 #status predicted <MAT>
F;22-374/Domain: extracellular #status predicted <EXT>
F;125-132/Region: acidic
F;169-230/Domain: immunoglobulin homology <IMM>
F;375-395/Domain: transmembrane #status predicted <TMM>
F;396-819/Domain: intracellular #status predicted <INT>
F;474-759/Domain: protein kinase homology <KIN>
F;482-490/Region: protein kinase ATP-binding motif
F;54-100,176-228,275-339/Disulfide bonds: #status predicted
F;76,116,225,238,262,294,315,328/Binding site: carbohydrate (Asn) (covalent) #status pre
F;512,529,621/Active site: Lys, Glu, Asp #status predicted
F;626,639/Binding site: magnesium (Asn, Asp) #status predicted
F;652/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicte

  Query Match             22.7%;  Score 389.5;  DB 1;  Length 819;
  Best Local Similarity   31.3%;  Pred. No. 1.5e-20;
  Matches   87;  Conservative   54;  Mismatches  118;  Indels   19;  Gaps    7;

Qy         23 AAARGPPKMADKVVPR-------QVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWS 75
              :|||  | : |: :|:         |  |  ::|:| :  |   :  | :||  :    :
Db         19 SAARPAPTLPDQALPKANIEVESHSAHPGDLLQLRCRLRDDVQSIN-WVRDGVQLPEN-N 76

Qy         76 RFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGG 135
              | |:  : ::|:    ||:|:| |   :  ||    |  | : |  |  |    :  |
Db         77 RTRITGEEVEVRDRVPEDSGLYACMTNSPSGS-ETTYFSVNVSDALPSAEDDDDEDDSSS 135

Qy        136 QEDPAS----QQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQAL 191
              :|  |      |   | :| | || ::: | |   :|: || : | | | : |:|: :
Db        136 EEKEADNTKPNQAVAPYWTYPEKMEKKLHAVPAAKTVKFKCPSGGTPNPTLRWLKNGKEF 195

Qy        192 TRPE----AAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPV 247
              :|:      : |   |:: : :: | | | ||| | |: |:|| ||::||::|:  :|:
Db        196 -KPDHRIGGYKVRYATWSIIMDSVVPSDKGNYTCIVENKYGSINHTYQLDVVERSPHRPI 254

Qy        248 LTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVE 285
              |    | | ||  |    | ||| || :| |||||  :|
Db        255 LQAGLPANKTVALGSNVEFVCKVYSDPQPHIQWLKHIE 292


RESULT 12
B54846
fibroblast growth factor receptor b precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 21-Jul-1995 #sequence_revision 28-Jul-1995 #text_change 16-Jul-1999
C;Accession: B54846
R;Takagi, Y.; Shrivastav, S.; Miki, T.; Sakaguchi, K.
J. Biol. Chem. 269, 23743-23749, 1994
A;Title: Molecular cloning and expression of the acidic fibroblast growth factor recepto
ility and covalent attachment of heparan sulfate glycosaminoglycan to the receptors.
A;Reference number: A54846; MUID:94375484; PMID:8089146
A;Accession: B54846
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-822 <TAK>
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; growth factor receptor
F;172-233/Domain: immunoglobulin homology <IMM>
F;480-765/Domain: protein kinase homology <KIN>
F;488-496/Region: protein kinase ATP-binding motif

  Query Match             22.7%;  Score 389.5;  DB 2;  Length 822;
  Best Local Similarity   31.3%;  Pred. No. 1.5e-20;
  Matches  100;  Conservative   56;  Mismatches  132;  Indels   31;  Gaps   10;

Qy         20 PPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRV 79
              ||        |: |  | |       | :: |:| ::     :  |||||  :| | :   |
Db         38 PPTKYQISQPE-ACVVAP------GESLELRCMLK--DAAVISWTKDG--VHLGPNNRTV 86

Qy         80 L-PQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED 138
              |  : |::|     |:|:| | |     | :: : : | | || | :    |:|     :
Db         87 LIGEYLQIKGATPRDSGLYACAAARTVDSETLYFMVNVTDAISSGDDEDDTDTSEDFVSE 146

Qy        139 PASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT---RPE 195
              | | | | :|    || :|: | |  ::|: :| | |:| | : |:|: :      |
Db        147 NRSNQRA-PYWTNTEKMEKRLHAVPAANTVKFRCPAGGNPTPTMRWLKNGKEFKQEHRIG 205

Qy        196 AAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVN 255
              : | : |:| :::: | | | ||| | |  |:|| || :||::|:  :|:|    | |
Db        206 GYKVRNQHWSLIMESVVPSDKGNYTCLVENEYGSINHTYHLDVVERSPHRPILQAGLPAN 265

Qy        256 TTVDFGGTTSFQCKVRSDVKPVIQWLKRVE-----YGAEG-------RHNSTIDVGGQKF 303
              :   ||    | ||| || :| |||:| ||     || :|       :|:    :
Db        266 ASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDGLPYLKVLKHSG---INSSNA 322

Qy        304 VVLPTGDVWSRPDGSYLNK 322
              ||   :|     | |: |
Db        323 EVLALFNVTEMDAGEYICK 341


RESULT 13
JC4058
fibroblast growth factor receptor-4 precursor - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Date: 29-Jun-1995 #sequence_revision 14-Jul-1995 #text_change 09-Jul-2004
C;Accession: JC4058
R;Shiozaki, C.; Tashiro, K.; Asano-Miyoshi, M.; Saigo, K.; Emori, Y.; Shiokawa, K.
Gene 152, 215-219, 1995
A;Title: Cloning of cDNA and genomic DNA encoding fibroblast growth factor rece ptor-4 of
A;Reference number: JC4058; MUID:95137391; PMID:7835703
A;Accession: JC4058
A;Molecule type: mRNA
A;Residues: 1-818 <SHI>
A;Cross-references: UNIPROT:Q91742; DDBJ:D31761; NID:g809527; PIDN:BAA06539.1; PID:g80952
C;Genetics:
A;Introns: 43/1; 125/1; 158/1; 213/1; 254/1; 318/1; 364/1; 429/1; 478/1; 556/1; 619/1; 68
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; protei
C;Keywords: ATP; growth factor receptor; transmembrane protein
F;1-26/Domain: signal sequence #status predicted <SIG>
F;27-818/Product: fibroblast growth factor receptor-4 #status predicted <MAT>
F;56-110/Domain: immunoglobulin homology <IM1>
F;132-137/Domain: acidic #status predicted <ADI>
F;177-238/Domain: immunoglobulin homology <IM2>
F;276-347/Domain: immunoglobulin homology <IM3>
F;382-402/Domain: transmembrane #status predicted <TMM>
F;477-762/Domain: protein kinase homology <KIN>
F;485-493/Region: protein kinase ATP-binding motif

  Query Match             22.5%;  Score 385.5;  DB 2;  Length 818;
  Best Local Similarity   29.5%;  Pred. No. 3e-20;
  Matches   98;  Conservative   55;  Mismatches  118;  Indels   61;  Gaps    9;

Qy         14 LLLGAFPPAAAARGPPKMAD--------KVVPRQVARLGRTVRLQCPVEGDPPPLTMWTK 65
              ||||     |  :   | :::         :|     :    |  :|| |             |
Db         18 LLLGVLLVATLSSCRPALSEDEANWKEPEVEEHLLLDPGNALRLFCD-----------TN 66

Qy         66 DGRTIHSGWSRFRVLPQG--------LKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVL 117
              :|:    : |:|| |         |:|   |  ||:|:|:|     | | :   ::: |:
Db         67 QSNSINWYREQDRLLPGGKIRMVGTVLEVSDVTYEDSGLYIC-VVRGTGKILRRFSISVV 125

Qy        118 DDISPGKESLGPDSSSGGQEDPASQ-------QWARPRFTQPSKMRRRVIARPVGSSVRL 170
```

```
                |  ::  | |    |   | :||  :          :  | :||| :| ::: | | |::|:
Db        126 DSLASGDEE---DDEDGRREDTTADINEEPVYFFQAPYWTQPHRMDKKLHAVPAGNTVKF 182

Qy        171 KCVASGHPRPDITWMKDDQALT---RPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRA 227
              :| | | | | | |:|: :      |      | : |:| :::: | | | ||| | ||
Db        183 RCPAGGSPLPTIRWLKNGREFRGEHRIGGIRLRHQHWSLVMESVVPSDRGNYTCVVENRV 242

Qy        228 GAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYG 287
              |::  || :||::|:  :|:|    | |||   |    | ||| || :| |||| :|
Db        243 GSLTYTYFLDVLERSSHRPILQAGLPANTTARVGSDVEFYCKVYSDAQPHIQWLKHIE-- 300

Qy        288 AEGRHNSTIDVGGQKF--------VVLPTGDV 311
                      | | .:|          || | |:
Db        301 ----------VNGSRFGPDDFPYVQVLKTADI 322


RESULT 14
S24108
protein-tyrosine kinase (EC 2.7.1.112) bek - chicken
N;Alternate names: receptor tyrosine kinase bek
C;Species: Gallus gallus (chicken)
C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004
C;Accession: S24108
R;Sato, M.; Kitazawa, T.; Iwai, T.; Seki, J.; Sakato, N.; Kato, J.; Takeya, T.
Oncogene 6, 1279-1283, 1991
A;Title: Isolation of chicken-bek and a related gene; identification of structural varia
A;Reference number: S24108; MUID:91319411; PMID:1650446
A;Accession: S24108
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-824 <SAT>
A;Cross-references: UNIPROT:Q90749; EMBL:X61992; NID:g63085; PIDN:CAA43965.1; PID:g63086
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; growth factor receptor; phosphotransferase; tyrosine-specific protein k
F;174-235/Domain: immunoglobulin homology <IMM>
F;482-767/Domain: protein kinase homology <KIN>
F;490-498/Region: protein kinase ATP-binding motif

  Query Match             22.3%;  Score 383.5;  DB 2;  Length 824;
  Best Local Similarity   30.0%;  Pred. No. 4.2e-20;
  Matches   91;  Conservative   57;  Mismatches  114;  Indels   41;  Gaps    9;

Qy         44 GRTVRLQCPVEGDPPPLTMWTKDG-------RTIHSGWSRFRVLPQGLKVKQVEREDAGV 96
              |  : |:| ::     :  |||||       ||:        :: : |::|     |:|:
Db         58 GEPLELRCQLK--DAVMISWTKDGVPLGPDNRTV--------IIGEYLQIKDASPRDSGL 107

Qy         97 YVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQED--PASQQWARPRFTQPSK 154
              | | |      | :: : : | | :| | :    :  : | ||    | |   | :|   |
Db        108 YACTAIRTLDSDTLYFIVNVTDALSSGDD----EDDNDGSEDFVNDSNQMRAPYWTHTDK 163

Qy        155 MRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALT---RPEAAEPRKKKWTLSLKNL 211
              | :|: | |  ::|: :| | |:| | : |:|: :      |    : | : |:| ::::
Db        164 MEKRLHAVPAANTVKFRCPAMGNPTPTMRWLKNGKEFKQEHRIGGYKVRNQHWSLIMESV 223

Qy        212 RPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVR 271
              | | | ||| | |: |:|| || :||::|:  :|:|    | | :   ||   | |||
Db        224 VPSDKGNYTCIVENQYGSINHTYHLDVVERSPHRPILQAGLPANASAVVGGDVEFVCKVY 283

Qy        272 SDVKPVIQWLKRVE-----YGAEG-------RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
              || :| |||:| ||     || :|       :|:    :      ||   :|     | |
Db        284 SDAQPHIQWIKHVERNGSKYGPDGLPYLQVLKHSG---INSSNAEVLTLYNVTEADAGEY 340

Qy        320 LNK 322
              : |
Db        341 ICK 343


RESULT 15
TVHUF4
fibroblast growth factor receptor 4 precursor - human
N;Alternate names: protein-tyrosine kinase tkf
N;Contains: protein-tyrosine kinase (EC 2.7.1.112)
C;Species: Homo sapiens (man)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 09-Jul-2004
C;Accession: S15345; A46615; A41598; D38269
R;Partanen, J.; Maekelae, T.P.; Eerola, E.; Korhonen, J.; Hirvonen, H.; Claesson-Welsh, I
EMBO J. 10, 1347-1354, 1991
A;Title: FGFR-4, a novel acidic fibroblast growth factor receptor with a distinct expres
A;Reference number: S15345; MUID:91224085; PMID:1709094
A;Accession: S15345
A;Molecule type: mRNA
A;Residues: 1-802 <PAR>
A;Cross-references: UNIPROT:P22455; EMBL:X57205; NID:g31371; PIDN:CAA40490.1; PID:g31372
A;Note: binds acidic but not basic fibroblast growth factor with high affinity
R;Ron, D.; Reich, R.; Chedid, M.; Lengel, C.; Cohen, O.E.; Chan, A.M.; Neufeld, G.; Miki,
J. Biol. Chem. 268, 5388-5394, 1993
A;Title: Fibroblast growth factor receptor 4 is a high affinity receptor for both acidic
A;Reference number: A46615; MUID:93194827; PMID:7680645
A;Accession: A46615
A;Status: nucleic acid sequence not shown; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 1-296,'D',298-802 <RON>
A;Experimental source: mammary epithelial cell line B5/589
A;Note: sequence extracted from NCBI backbone (NCBIP:127650)
A;Note: binds acidic and basic fibroblast growth factors with high affinity
R;Holtrich, U.; Braeuninger, A.; Strebhardt, K.; Ruebsamen-Waigmann, H.
Proc. Natl. Acad. Sci. U.S.A. 88, 10411-10415, 1991
A;Title: Two additional protein-tyrosine kinases expressed in human lung: fourth member c
A;Reference number: S19025; MUID:92073297; PMID:1720539
A;Accession: A41598
A;Status: nucleic acid sequence not shown; not compared with conceptual translation
A;Molecule type: mRNA
A;Residues: 399-534,'M',536-799,'SG',800-802 <HOL>
A;Experimental source: lung
R;Partanen, J.; Maekelae, T.P.; Alitalo, R.; Lehvaeslaiho, H.; Alitalo, K.
Proc. Natl. Acad. Sci. U.S.A. 87, 8913-8917, 1990
A;Title: Putative tyrosine kinases expressed in K-562 human leukemia cells.
A;Reference number: A38268; MUID:91062389; PMID:2247464
A;Accession: D38269
A;Molecule type: mRNA
A;Residues: 614-670 <PA2>
A;Cross-references: GB:M37781
A;Experimental source: K-562 leukemia cell line
C;Genetics:
A;Gene: GDB:FGFR4
A;Cross-references: GDB:127929; OMIM:134935
A;Map position: 5q33.2-5qter
C;Function:
A;Description: receptor mediating effects of fibroblast growth factor
A;Note: expressed in normal lung; expressed in some carcinomas
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prote
C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor receptor;
F;1-24/Domain: signal sequence #status predicted <SIG>
F;25-802/Product: fibroblast growth factor receptor 4 #status predicted <MAT>
F;25-369/Domain: extracellular #status predicted <EXT>
F;50-103/Domain: immunoglobulin homology <IM1>
F;165-226/Domain: immunoglobulin homology <IM2>
F;264-335/Domain: immunoglobulin homology <IM3>
F;370-390/Domain: transmembrane #status predicted <TMM>
F;391-802/Domain: intracellular #status predicted <INT>
F;465-750/Domain: protein kinase homology <KIN>
F;473-481/Region: protein kinase ATP-binding motif
F;57-101,172-224,271-333/Disulfide bonds: #status predicted
F;112,258,290,311,322/Binding site: carbohydrate (Asn) (covalent) #status predicted
F;503,520,612/Active site: Lys, Glu, Asp #status predicted
F;617,630/Binding site: magnesium (Asn, Asp) #status predicted
F;643/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted

  Query Match             22.3%;  Score 382.5;  DB 1;  Length 802;
  Best Local Similarity   31.3%;  Pred. No. 4.8e-20;
  Matches  109;  Conservative   50;  Mismatches  136;  Indels   53;  Gaps   12;

Qy          6 LLLLLLPPLLLGAFPPAAAARG------PPKMADKVVPRQ---VARLGRTVRLQCPVEGD 56
              ||| ||  ||     ||  :          | :|  :  ::      ||: ||| |   |
Db          3 LLLALLGVLLSVPGPPVLSLEASEEVELEPCLAPSLEQQEQELTVALGQPVRLCC---GR 59
```

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Qy          57 PPPLTMWTKDGRTIHSGWSRFRVLPQG--------LKVKQVEREDAGVYVCKATNGFGSL 108
               |  |:|         .  |: | |          |::     ||||  |:|  |     ||:
Db          60 AERGGHWYKEGS---------RLAPAGRVRGWRGRLEIASFLPEDAGRYLCLAR---GSM 107

Qy         109 SV--NYTLVVLDDISPGKESLGPDSSSGGQEDPA---SQQWARPRFTQPSKMRRRVIARP 163
                |  | ||:  | ::   :   |.|       ||:   |      | :| | :| ::: | |
Db         108 IVLQNLTLITGDSLTSSNDDEDPKS----HRDPSNRHSYPQQAPYWTHPQRMEKKLHAVP 163

Qy         164 VGSSVRLKCVASGHPRPDITWMKDDQAL---TRPEAAEPRKKKWTLSLKNLRPEDSGKYT 220
                |::|: :| |:|:| | | |:|| ||     |      | : |:| :::: | | | ||
Db         164 AGNTVKFRCPAAGNPTPTIRWLKDGQAFHGENRIGGIRLRHQHWSLVMESVVPSDRGTYT 223

Qy         221 CRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQW 280
               | | |  |:|   | :||::|:  :|:|    | |||   |      ||| || :| |||
Db         224 CLVENAVGSIRYNYLLDVLERSPHRPILQAGLPANTTAVVGSDVELLCKVYSDAQPHIQW 283

Qy         281 LKRV-----EYGAEG----RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
               || :      :|| |    :   | |:   :  ||   :| :   | |
Db         284 LKHIVINGSSFGAVGFPYVQVLKTADINSSEVEVLYLRNVSAEDAGEY 331
```

Search completed: February 2, 2005, 18:26:28
Job time : 22 secs

GenCore version 5.1.6

OM protein - protein search, using sw model

Run on: February 2, 2005, 18:16:50 ; Search time 194 Seconds
(without alignments)
960.935 Million cell updates/sec

Title: US-10-613-413B-8
Perfect score: 1717
Sequence: 1 MTPSPLLLLLLPPLLLGAFP.....VLPTGDVWSRPDGSYLNKPL 324

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1825181 seqs, 575374646 residues

Total number of hits satisfying chosen parameters: 1825181

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt_02:*
1: uniprot_sprot:*
2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1707	99.4	497	2	Q9BXN7	Q9bxn7 homo sapien
2	1707	99.4	504	2	Q8N441	Q8n441 homo sapien
3	1707	99.4	504	2	Q9H4D7	Q9h4d7 homo sapien
4	1707	99.4	504	2	AAQ88670	Aaq88670 homo sapi
5	1561	90.9	529	2	Q7TQM3	Q7tqm3 rattus norv
6	1560	90.9	529	2	Q91V87	Q91v87 mus musculu
7	1560	90.9	529	2	AAH58745	Aah58745 mus muscu
8	1553	90.4	440	2	Q6ZMD4	Q6zmd4 homo sapien
9	1553	90.4	440	2	BAD18794	Bad18794 homo sapi
10	1507	87.8	509	2	Q920C2	Q920c2 mus musculu
11	1306	76.1	487	2	Q7T2H2	Q7t2h2 gallus gall
12	1193.5	69.5	483	2	Q7SX76	Q7sx76 brachydanio
13	1035.5	60.3	438	2	Q920C3	Q920c3 mus musculu
14	894	52.1	350	2	Q6PJN1	Q6pjn1 homo sapien
15	894	52.1	350	2	AAH13955	Aah13955 homo sapi
16	425.5	24.8	812	1	FGR1_XENLA	P22182 xenopus lae
17	424.5	24.7	800	2	Q9I8X3	Q9i8x3 brachydanio
18	424.5	24.7	814	2	Q91897	Q91897 xenopus lae
19	423.5	24.7	810	2	Q9PS96	Q9ps96 xenopus lae
20	422	24.6	822	2	Q91288	Q91288 pleurodeles
21	415.5	24.2	802	2	O42127	O42127 xenopus lae
22	413	24.1	827	2	Q6GNS5	Q6gns5 xenopus lae
23	409	23.8	800	2	Q99052	Q99052 mus musculu
24	408.5	23.8	800	2	Q7TSI8	Q7tsi8 mus musculu
25	408.5	23.8	801	1	FGR3_MOUSE	Q61851 mus musculu
26	399.5	23.3	796	2	Q91287	Q91287 pleurodeles
27	399.5	23.3	802	2	Q95M13	Q95m13 bos taurus
28	398	23.2	370	2	Q800Y8	Q800y8 brachydanio
29	398	23.2	800	2	Q9JHX9	Q9jhx9 rattus norv
30	394.5	23.0	771	2	Q8NI16	Q8ni16 homo sapien
31	393.5	22.9	806	1	CEK2_CHICK	P18460 gallus gall
32	393	22.9	806	1	FGR3_HUMAN	P22607 homo sapien
33	392.5	22.9	446	2	Q63236	Q63236 rattus norv
34	392.5	22.9	819	1	FGR1_CHICK	P21804 gallus gall
35	392	22.8	692	2	Q800Y9	Q800y9 brachydanio
36	392	22.8	756	2	Q800Z0	Q800z0 brachydanio
37	392	22.8	804	2	Q800Z1	Q800z1 brachydanio
38	392	22.8	806	2	Q90Z00	Q90z00 brachydanio
39	389.5	22.7	769	2	Q8NI15	Q8ni15 homo sapien
40	389.5	22.7	822	2	Q9QVV7	Q9qvv7 rattus sp.
41	388.5	22.6	446	2	Q63237	Q63237 rattus norv
42	387.5	22.6	807	2	Q6DD66	Q6dd66 xenopus lae
43	387.5	22.6	818	2	Q9PSV9	Q9psv9 xenopus lae
44	387.5	22.6	828	2	Q9DGK3	Q9dgk3 xenopus lae
45	387	22.5	815	2	Q805B9	Q805b9 brachydanio

ALIGNMENTS

RESULT 1
Q9BXN7
ID Q9BXN7 PRELIMINARY; PRT; 497 AA.
AC Q9BXN7;
DT 01-JUN-2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE FGF homologous factor receptor.
GN Name=FHFR;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA Aggarwal S., Xie M.-H., Foster J., Frantz G., Stinson J., Corpuz R.T.,
RA Simmons L., Hillan K., Yansura D.G., Vandlen R.L., Goddard A.D.,
RA Gurney A.L.;
RL Submitted (OCT-2000) to the EMBL/GenBank/DDBJ databases.
DR EMBL; AF312678; AAK15273.1; -.
DR HSSP; P11362; 1EVT.
DR GO; GO:0004872; F:receptor activity; IEA.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 3.
DR SMART; SM00408; IGc2; 2.
DR PROSITE; PS50835; IG_LIKE; 3.
KW Receptor.
SQ SEQUENCE 497 AA; 53757 MW; 57301F4F36357360 CRC64;

Query Match 99.4%; Score 1707; DB 2; Length 497;
Best Local Similarity 99.7%; Pred. No. 9.1e-124;
Matches 323; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||||||| |
Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 2
Q8N441
ID   Q8N441      PRELIMINARY;      PRT;   504 AA.
AC   Q8N441;
DT   01-OCT-2002 (TrEMBLrel. 22, Created)
DT   01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT   01-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Fibroblast growth factor receptor-like 1, (FGFRL1).
GN   Name=FGFRL1; ORFNames=UNQ480;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Brain;
RX   MEDLINE=22388257; PubMed=12477932;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Brain;
RA   Strausberg R.;
RL   Submitted (AUG-2002) to the EMBL/GenBank/DDBJ databases.
RN   [3]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=22887296; PubMed=12975309;
RA   Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
RA   Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA   Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
RA   Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
RA   Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
RA   Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA   Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D.,
RA   Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
RA   Godowski P.;
RT   "The secreted protein discovery initiative (SPDI), a large-scale
RT   effort to identify novel human secreted and transmembrane proteins: a
RT   bioinformatics assessment.";
RL   Genome Res. 13:2265-2270(2003).
DR   EMBL; BC036769; AAH36769.1; -.
DR   EMBL; AY358303; AAQ88670.1; -.
DR   HSSP; P11362; 1EVT.
DR   Genew; HGNC:3693; FGFRL1.
DR   GO; GO:0004872; F:receptor activity; IEA.
DR   InterPro; IPR007110; Ig-like.
DR   Pfam; PF00047; ig; 3.
DR   PROSITE; PS50835; IG_LIKE; 3.
KW   Receptor.
SQ   SEQUENCE   504 AA;  54536 MW;  56E35E57D5FC141B CRC64;

  Query Match          99.4%;  Score 1707;  DB 2;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 9.2e-124;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
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Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324


RESULT 3
Q9H4D7
ID   Q9H4D7      PRELIMINARY;      PRT;   504 AA.
AC   Q9H4D7;
DT   01-MAR-2001 (TrEMBLrel. 16, Created)
DT   01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   FGFR-like protein precursor (Fibroblast growth factor receptor
DE   5).
GN   Name=FGFRL1; Synonyms=FGFR5;
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Cartilage;
RX   PubMed=11031111;
RA   Wiedemann M., Trueb B.;
RT   "Characterization of a novel protein (FGFRL1) from human cartilage
RT   related to FGF receptors.";
RL   Genomics 69:275-279(2000).
RN   [2]
RP   SEQUENCE FROM N.A.
RX   MEDLINE=21167383; PubMed=11267671;
RA   Kim I., Moon S.O., Yu K.H., Kim U.H., Koh G.Y.;
RT   "A novel fibroblast growth factor receptor-5 preferentially expressed
RT   in the pancreas.";
RL   Biochim. Biophys. Acta 1518:152-156(2001).
DR   EMBL; AJ277437; CAC14171.1; -.
DR   EMBL; AF279689; AAK26742.1; -.
DR   HSSP; P11362; 1EVT.
DR   GO; GO:0016021; C:integral to membrane; NAS.
DR   GO; GO:0005007; F:fibroblast growth factor receptor activity; NAS.
DR   GO; GO:0001558; P:regulation of cell growth; NAS.
DR   InterPro; IPR007110; Ig-like.
DR   InterPro; IPR003598; Ig_c2.
DR   Pfam; PF00047; ig; 3.
DR   SMART; SM00408; IGc2; 2.
DR   PROSITE; PS50835; IG_LIKE; 3.
KW   Receptor; Signal.
FT   SIGNAL        1     17       Potential.
SQ   SEQUENCE   504 AA;  54567 MW;  16382E57D4276485 CRC64;
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  Query Match             99.4%;  Score 1707;  DB 2;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 9.2e-124;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 4
AAQ88670
ID AAQ88670 PRELIMINARY; PRT; 504 AA.
AC AAQ88670;
DT 02-MAR-2004 (TrEMBLrel. 27, Created)
DT 02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
DT 02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
DE FGFRL1.
GN UNQ480.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primata; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RX PubMed=12975309;
RA Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,
RA Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.,
RA Eaton D., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,
RA Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,
RA Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,
RA Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,
RA Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D.,
RA Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
RA Godowski P.;
RT "The Secreted Protein Discovery Initiative (SPDI), a Large-Scale
RT Effort to Identify Novel Human Secreted and Transmembrane Proteins: A
RT Bioinformatics Assessment.";
RL Genome Res. 13:2265-2270(2003).
DR EMBL; AY358303; AAQ88670.1; -.
SQ SEQUENCE 504 AA; 54536 MW; 56E35E57D5FC141B CRC64;

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  Query Match             99.4%;  Score 1707;  DB 2;  Length 504;
  Best Local Similarity   99.7%;  Pred. No. 9.2e-124;
  Matches  323;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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Db        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
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Db        301 QKFVVLPTGDVWSRPDGSYLNKLL 324
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RESULT 5
Q7TQM3
ID Q7TQM3 PRELIMINARY; PRT; 529 AA.
AC Q7TQM3;
DT 01-OCT-2003 (TrEMBLrel. 25, Created)
DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Fibroblast growth factor receptor-like protein precursor.
GN Name=fgfrl1;
OS Rattus norvegicus (Rat).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
OX NCBI_TaxID=10116;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=Sprague-Dawley;
RX PubMed=12813049;
RA Trueb B., Zhuang L., Taeschler S., Wiedemann M.;
RT "Characterization of FGFRL1, a Novel FGF receptor preferentially
RT expressed in skeletal tissues.";
RL J. Biol. Chem. 278:33857-33865(2003).
DR EMBL; AJ536020; CAD59914.1; -.
DR GO; GO:0004872; F:receptor activity; IEA.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 3.
DR SMART; SM00408; IGc2; 3.
DR PROSITE; PS50835; IG_LIKE; 3.
KW Receptor; Signal.
FT SIGNAL 1 20 Potential.
FT CHAIN 21 529 fibroblast growth factor receptor-like
FT protein.
SQ SEQUENCE 529 AA; 57143 MW; 23D67B419335FFFC CRC64;

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  Query Match             90.9%;  Score 1561;  DB 2;  Length 529;
  Best Local Similarity   91.4%;  Pred. No. 2e-112;
  Matches  296;  Conservative    9;  Mismatches   15;  Indels    4;  Gaps    1;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    ||||| | | ||||||:||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
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Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||: ||  |||||||| ||||||||||||||||||||||||||||||||||||||
Db        117 SPGKENPGPGGSSGGQEDPVSQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||| ||:| ||||||||||||:|||||||||||||||||||||||||||
Db        177 DIMWMKDDQTLTRLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNRAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
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                ||||||||||||||||||||||||||||||||||||||||||||||||:||||||||||||
Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
                  ||||||||||||||||||||| |
Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 6
Q91V87
ID   Q91V87      PRELIMINARY;      PRT;   529 AA.
AC   Q91V87;
DT   01-DEC-2001 (TrEMBLrel. 19, Created)
DT   01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT   01-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   Fibroblast growth factor receptor 5 beta precursor (FGF receptor-like
DE   protein 1 precursor) (Fibroblast growth factor receptor-like 1).
GN   Name=Fgfrl1; Synonyms=FGFRL1, Fgfr5;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB/cJ;
RX   MEDLINE=21311632; PubMed=11418238;
RA   Sleeman M., Fraser J., McDonald M., Yuan S., White D., Grandison P.,
RA   Kumble K., Watson J.D., Murison J.G.;
RT   "Identification of a new fibroblast growth factor receptor, FGFR5.";
RL   Gene 271:171-182(2001).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB/cJ;
RA   Sleeman M.A., Fraser J.K., Watson J.D., Kumble K.D., Murison J.G.;
RL   Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.
RN   [3]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Cartilage;
RX   MEDLINE=21450304; PubMed=11566361;
RA   Wiedemann M., Trueb B.;
RT   "The mouse Fgfrl1 gene coding for a novel FGF receptor-like protein.";
RL   Biochim. Biophys. Acta 1520:247-250(2001).
RN   [4]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB /c; TISSUE=Liver;
RX   PubMed=11031111;
RA   Wiedemann M., Trueb B.;
RT   "Characterization of a novel protein (FGFRL1) from human cartilage
RT   related to FGF receptors.";
RL   Genomics 69:275-279(2000).
RN   [5]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB /c; TISSUE=Liver;
RA   Wiedemann M., Trueb B.;
RL   Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.
RN   [6]
RP   SEQUENCE FROM N.A.
RC   STRAIN=FVB/N-3;
RC   TISSUE=Mammary tumor. MMTV-LTR/INT3 model. 5 month old mouse. Taken by
RC   biopsy.;
RX   MEDLINE=22388257; PubMed=12477932;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [7]
RP   SEQUENCE FROM N.A.
RC   STRAIN=FVB/N-3;
RC   TISSUE=Mammary tumor. MMTV-LTR/INT3 model. 5 month old mouse. Taken by
RC   biopsy.;
RA   Strausberg R.;
RL   Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AF321300; AAL06295.1; -.
DR   EMBL; AJ293947; CAC82376.1; -.
DR   EMBL; AJ308490; CAC83768.1; -.
DR   EMBL; BC058745; AAH58745.1; -.
DR   HSSP; P11362; 1EVT.
DR   MGD; MGI:2150920; Fgfrl1.
DR   GO; GO:0005886; C:plasma membrane; IDA.
DR   GO; GO:0005515; F:protein binding; IPI.
DR   GO; GO:0008285; P:negative regulation of cell proliferation; IDA.
DR   InterPro; IPR007110; Ig-like.
DR   Pfam; PF00047; ig; 3.
DR   PROSITE; PS50835; IG_LIKE; 3.
KW   Receptor; Signal.
FT   SIGNAL       1     20       Potential.
FT   CHAIN       21    529       fibroblast growth factor receptor 5 beta.
SQ   SEQUENCE   529 AA;  57013 MW;  6CE76A75BF2498DD CRC64;

  Query Match             90.9%;  Score 1560;  DB 2;  Length 529;
  Best Local Similarity   91.4%;  Pred. No. 2.4e-112;
  Matches  296;  Conservative    9;  Mismatches   15;  Indels    4;  Gaps    1;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    ||||| | | ||||||:||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||:|| ||||||||||||||||||:::|||
Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||
Db        117 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| ||||||||||:||||||||||:||||||||||||||||||
Db        177 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||||||:||||||||||||
Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||| |
Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 7
AAH58745
ID   AAH58745    PRELIMINARY;      PRT;   529 AA.
AC   AAH58745;
DT   02-MAR-2004 (TrEMBLrel. 27, Created)
DT   02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
DT   02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
DE   Fibroblast growth factor receptor-like 1.
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
```

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RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=FVB/N-3; TISSUE=Mammary tumor;
RX   MEDLINE=22388257; PubMed=12477932;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=FVB/N-3; TISSUE=Mammary tumor;
RA   Strausberg R.;
RL   Submitted (SEP-2003) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; BC058745; AAH58745.1; -.
KW   Receptor.
SQ   SEQUENCE   529 AA;  57013 MW;  6CE76A75BF2498DD CRC64;

  Query Match             90.9%;  Score 1560;  DB 2;  Length 529;
  Best Local Similarity   91.4%;  Pred. No. 2.4e-112;
  Matches  296;  Conservative    9;  Mismatches   15;  Indels    4;  Gaps    1;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    ||||| | | ||||||:||||||||||||||||||||||||||||||
Db          1 MTRSPALLL----LLLGALPSAEAARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPL 56

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
              ||||||||||||||||||||||||||:|| ||||||||||||||||||||||:::|||
Db         57 TMWTKDGRTIHSGWSRFRVLPQGLKVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDI 116

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              |||||| ||  ||||||||||||||||||||||||||||||||||||||||||||||||
Db        117 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 176

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || |||||| ||  ||:| |||||||||||:||||||||||:|||||||||||||||
Db        177 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 236

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||:|||||||||||||
Db        237 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 296

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||| |
Db        297 QKFVVLPTGDVWSRPDGSYLNKLL 320


RESULT 8
Q6ZMD4
ID   Q6ZMD4      PRELIMINARY;      PRT;   440 AA.
AC   Q6ZMD4;
DT   05-JUL-2004 (TrEMBLrel. 27, Created)
DT   05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT   05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE   Hypothetical protein FLJ23990.
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Kawabata A., Hikiji T., Kobatake N., Inagaki H., Ikema Y., Okamoto S.,
RA   Okitani R., Ota T., Suzuki Y., Obayashi M., Nishi T., Shibahara T.,
RA   Tanaka T., Nakamura Y., Isogai T., Sugano S.;
RL   Submitted (APR-2004) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AK172829; BAD18794.1; -.
DR   GO; GO:0004872; F:receptor activity; IEA.
DR   InterPro; IPR003599; Ig.
DR   InterPro; IPR007110; Ig-like.
DR   InterPro; IPR003598; Ig_c2.
DR   Pfam; PF00047; ig; 3.
DR   SMART; SM00409; IG; 3.
DR   SMART; SM00408; IGc2; 3.
DR   PROSITE; PS50835; IG_LIKE; 3.
KW   Receptor.
SQ   SEQUENCE   440 AA;  47587 MW;  2B6E0C2285AC3B59 CRC64;

  Query Match             90.4%;  Score 1553;  DB 2;  Length 440;
  Best Local Similarity   99.7%;  Pred. No. 6.6e-112;
  Matches  293;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          31 MADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVE 90
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db           1 MADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVE 60

Qy          91 REDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFT 150
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          61 REDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFT 120

Qy         151 QPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKN 210
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         121 QPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKN 180

Qy         211 LRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKV 270
               ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         181 LRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKV 240

Qy         271 RSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
               ||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db         241 RSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKLL 294


RESULT 9
BAD18794
ID   BAD18794    PRELIMINARY;      PRT;   440 AA.
AC   BAD18794;
DT   12-MAY-2004 (TrEMBLrel. 27, Created)
DT   12-MAY-2004 (TrEMBLrel. 27, Last sequence update)
DT   12-MAY-2004 (TrEMBLrel. 27, Last annotation update)
DE   CDNA FLJ23990 fis, clone HRC08053, highly similar to Homo sapiens
DE   fibroblast growth factor receptor-like 1 (FGFRL1).
OS   Homo sapiens (Human).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX   NCBI_TaxID=9606;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Kawabata A., Hikiji T., Kobatake N., Inagaki H., Ikema Y., Okamoto S.,
RA   Okitani R., Ota T., Suzuki Y., Obayashi M., Nishi T., Shibahara T.,
RA   Tanaka T., Nakamura Y., Isogai T., Sugano S.;
RT   "NEDO human cDNA sequencing project.";
RL   Submitted (APR-2004) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AK172829; BAD18794.1; -.
KW   Receptor.
SQ   SEQUENCE   440 AA;  47587 MW;  2B6E0C2285AC3B59 CRC64;

  Query Match             90.4%;  Score 1553;  DB 2;  Length 440;
  Best Local Similarity   99.7%;  Pred. No. 6.6e-112;
  Matches  293;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy          31 MADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVE 90
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Db          1 MADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGLKVKQVE 60

Qy         91 REDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFT 150
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 REDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQWARPRFT 120

Qy        151 QPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKN 210
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 QPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKN 180

Qy        211 LRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKV 270
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 LRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKV 240

Qy        271 RSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        241 RSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKLL 294
```

RESULT 10
Q920C2
ID Q920C2 PRELIMINARY; PRT; 509 AA.
AC Q920C2;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Fibroblast growth factor receptor 5 beta/gamma (Fragment).
GN Name=Fgfrl1; Synonyms=Fgfr5;
OS Mus musculus (Mouse).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX NCBI_TaxID=10090;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=BALB/cJ;
RX MEDLINE=21311632; PubMed=11418238;
RA Sleeman M., Fraser J., McDonald M., Yuan S., White D., Grandison P.,
RA Kumble K., Watson J.D., Murison J.G.;
RT "Identification of a new fibroblast growth factor receptor, FGFR5.";
RL Gene 271:171-182(2001).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=BALB/cJ;
RA Sleeman M.A., Fraser J.K., McDonald M., Yuan S., White D.,
RA Grandison P., Kumble K.D., Watson J.D., Murison J.G.;
RL Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.
DR EMBL; AF321302; AAL06297.1; -.
DR HSSP; P11362; 1EVT.
DR MGD; MGI:2150920; Fgfrl1.
DR GO; GO:0005886; C:plasma membrane; IDA.
DR GO; GO:0005515; F:protein binding; IPI.
DR GO; GO:0008285; P:negative regulation of cell proliferation; IDA.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 3.
DR SMART; SM00408; IGc2; 3.
DR PROSITE; PS50835; IG_LIKE; 3.
KW Receptor.
FT NON_TER 1 1
FT CHAIN <1 509 fibroblast growth factor receptor 5 beta.
FT CHAIN <1 509 fibroblast growth factor receptor 5
FT gamma.
SQ SEQUENCE 509 AA; 54994 MW; 392A7D9568899C15 CRC64;

Query Match 87.8%; Score 1507; DB 2; Length 509;
Best Local Similarity 93.7%; Pred. No. 2.9e-108;
Matches 281; Conservative 9; Mismatches 10; Indels 0; Gaps 0;

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Qy         25 ARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGL 84
              |||||:||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 ARGPPRMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGL 60
```

```
Qy         85 KVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQW 144
              |||:|| |||||||||||||||||||||:::|||||||| ||  ||||||||||||||
Db         61 KVKEVEAEDAGVYVCKATNGFGSLSVNYTLIIMDDISPGKESPGPGGSSGGQEDPASQQW 120

Qy        145 ARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKW 204
              |||||||||||||||||||||||||||||||||||| |||||| ||  ||:| |||||
Db        121 ARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDIMWMKDDQTLTHLEASEHRKKKW 180

Qy        205 TLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTT 264
              |||||||:|||||||||||||:|||||||||||||||||||||||||||||||||||||
Db        181 TLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTT 240

Qy        265 SFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||:|||||||||||||||||||||||||||||||||||||| |
Db        241 SFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKLL 300
```

RESULT 11
Q7T2H2
ID Q7T2H2 PRELIMINARY; PRT; 487 AA.
AC Q7T2H2;
DT 01-OCT-2003 (TrEMBLrel. 25, Created)
DT 01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE Fibroblast growth factor receptor-like protein precursor.
GN Name=FGFRL1;
OS Gallus gallus (Chicken).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
OC Gallus.
OX NCBI_TaxID=9031;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Cartilage;
RX PubMed=12813049;
RA Trueb B., Zhuang L., Taeschler S., Wiedemann M.;
RT "Characterization of FGFRL1, a Novel FGF receptor preferentially
RT expressed in skeletal tissues.";
RL J. Biol. Chem. 278:33857-33865(2003).
DR EMBL; AJ535114; CAD59380.1; -.
DR GO; GO:0004872; F:receptor activity; IEA.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 3.
DR SMART; SM00408; IGc2; 3.
DR PROSITE; PS50835; IG_LIKE; 3.
KW Receptor; Signal.
FT SIGNAL 1 18 Potential.
FT CHAIN 19 487 fibroblast growth factor receptor-like
FT protein.
SQ SEQUENCE 487 AA; 54099 MW; FFD0132AD917FF94 CRC64;

Query Match 76.1%; Score 1306; DB 2; Length 487;
Best Local Similarity 75.8%; Pred. No. 9.9e-93;
Matches 238; Conservative 36; Mismatches 40; Indels 0; Gaps 0;

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Qy          9 LLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGR 68
              | |  || |    : :|||||::||||: ||  |||||::| |||||||||||| ||||
Db          3 LQLALLLAGIVALSDSARGPPRIADKVIHRQSVRLGRTIKLLCPVEGDPPPLTMWMKDGR 62

Qy         69 TIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLG 128
              ||||||:|||:| ||||:|:|| |||| |:||||||||| :|||||:|:|| | || |
Db         63 TIHSGWTRFRILQQGLKIKEVESEDAGTYICKATNGFGSTNVNYTLIVIDDTSSGKNSQT 122

Qy        129 PDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDD 188
              |: |:|  || : :|||:||||||:||||||||||||||:||||||||:|||||||:||:
Db        123 PEGSNGEYEDHSGKQWAQPRFTQPAKMRRRVIARPVGSSIRLKCVASGNPRPDITWLKDN 182

Qy        189 QALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVL 248
              : |   |  | :||||||:||||:||||||||||| |: | |||||||:|||||||:|
Db        183 KPLMPHEIGENKKKKWTLNLKNLKPEDSGKYTCRVFNKVGEINATYKVEVIQRTRSKPIL 242

Qy        249 TGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPT 308
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                ||||||||||:|||||||||||||||||||||||||||| | ::|||||||||||||||||
Db        243 TGTHPVNTTVDYGGTTSFQCKVRSDVKPVIQWLKRVEYGTESKYNSTIDVGGQKFVVLPT 302

Qy        309 GDVWSRPDGSYLNK 322
              |:||||||||||||
Db        303 GEVWSRPDGSYLNK 316


RESULT 12
Q7SX76
ID   Q7SX76      PRELIMINARY;      PRT;   483 AA.
AC   Q7SX76;
DT   01-OCT-2003 (TrEMBLrel. 25, Created)
DT   01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
DT   01-OCT-2004 (TrEMBLrel. 28, Last annotation update)
DE   FGF receptor-like protein precursor (Fibroblast growth factor
DE   receptor-like 1).
GN   Name=fgfrl1;
OS   Brachydanio rerio (Zebrafish) (Danio rerio).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;
OC   Cyprinidae; Danio.
OX   NCBI_TaxID=7955;
RN   [1]
RP   SEQUENCE FROM N.A.
RA   Trueb B., Zhuang L., Taeschler S., Wiedemann M.;
RT   "Characterization of FGFRL1, a novel FGF receptor preferentially
RT   expressed in skeletal tissues.";
RL   J. Biol. Chem. 278:0-0(2003).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Kidney;
RX   MEDLINE=22388257; PubMed=12477932;
RA   Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA   Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA   Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA   Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA   Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA   Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA   Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA   Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA   Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA   Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA   Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA   Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA   Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA   Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA   Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA   Jones S.J., Marra M.A.;
RT   "Generation and initial analysis of more than 15,000 full-length human
RT   and mouse cDNA sequences.";
RL   Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN   [3]
RP   SEQUENCE FROM N.A.
RC   TISSUE=Kidney;
RA   Strausberg R.;
RL   Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AJ574916; CAE00573.1; -.
DR   EMBL; BC053245; AAH53245.1; -.
DR   ZFIN; ZDB-GENE-040128-2; fgfrl1.
DR   GO; GO:0004872; F:receptor activity; IEA.
DR   InterPro; IPR007110; Ig-like.
DR   InterPro; IPR003598; Ig_c2.
DR   Pfam; PF00047; ig; 3.
DR   SMART; SM00408; IGc2; 3.
DR   PROSITE; PS50835; IG_LIKE; 3.
KW   Receptor; Signal.
FT   SIGNAL       1     19       Potential.
FT   CHAIN       20    483       FGF receptor-like protein.
SQ   SEQUENCE   483 AA;  53937 MW;  3C2B34E5D57C332C CRC64;

  Query Match             69.5%;  Score 1193.5;  DB 2;  Length 483;
  Best Local Similarity   73.0%;  Pred. No. 5.1e-84;
  Matches  219;  Conservative   33;  Mismatches   47;  Indels    1;  Gaps    1;

Qy         25 ARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGRTIHSGWSRFRVLPQGL 84
              |||||::|:|:  ||  |:|||::|||||||||||| |||||| ||||| ||||| | |
Db         20 ARGPPRVAEKIAHRQTVRIGRTMKLQCPVEGDPPPLIMWTKDGRNIHSGWMRFRVLQQAL 79

Qy         85 KVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDISPGKESLGPDSSSGGQEDPASQQW 144
              ::|:|| :||| ::|||||||||:::||||:|:|| | |:|   |   :   | : :
Db         80 RIKEVEADDAGTFICKATNGFGSVNININYTLIVIDDSSAGREGARPAGETEYSTD-LTGKL 138

Qy        145 ARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKW 204
              |||||||:|||:|||||||||||||| |||:||||| |:|| : ||  |  | |||||
Db        139 VRPRFTQPAKMRKRVIARPVGSSVRLKCTASGNPRPDIVWLKDSRPLTPEEVGEGRKKKW 198

Qy        205 TLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTT 264
              ||||||| || |||||| |||||| |||||||:||||| |||:||||||||||:||||
Db        199 TLSLKNLTPEHSGKYTCHVSNRAGEINATYKVEVIQRTNSKPILTGTHPVNTTVDYGGTT 258

Qy        265 SFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||| | ||::||||:||   |||||||||||||||||||| |
Db        259 SFQCKVRSDVKPVIQWLKRVEPGGEGKYNSTIEVGDHHFVVLPTGDVWSRPDGSYLNKLL 318


RESULT 13
Q920C3
ID   Q920C3      PRELIMINARY;      PRT;   438 AA.
AC   Q920C3;
DT   01-DEC-2001 (TrEMBLrel. 19, Created)
DT   01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT   01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DE   Fibroblast growth factor receptor 5 gamma precursor.
GN   Name=Fgfrl1; Synonyms=Fgfr5;
OS   Mus musculus (Mouse).
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC   Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
OX   NCBI_TaxID=10090;
RN   [1]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB/cJ;
RX   MEDLINE=21311632; PubMed=11418238;
RA   Sleeman M., Fraser J., McDonald M., Yuan S., White D., Grandison P.,
RA   Kumble K., Watson J.D., Murison J.G.;
RT   "Identification of a new fibroblast growth factor receptor, FGFR5.";
RL   Gene 271:171-182(2001).
RN   [2]
RP   SEQUENCE FROM N.A.
RC   STRAIN=BALB/cJ;
RA   Sleeman M.A., Fraser J.K., Watson J.D., Kumble K.D., Murison J.G.;
RL   Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.
DR   EMBL; AF321301; AAL06296.1; -.
DR   HSSP; P11362; 1EVT.
DR   MGD; MGI:2150920; Fgfrl1.
DR   GO; GO:0005886; C:plasma membrane; IDA.
DR   GO; GO:0005515; F:protein binding; IPI.
DR   GO; GO:0008285; P:negative regulation of cell proliferation; IDA.
DR   InterPro; IPR007110; Ig-like.
DR   InterPro; IPR003598; Ig_c2.
DR   Pfam; PF00047; ig; 2.
DR   SMART; SM00408; IGc2; 2.
DR   PROSITE; PS50835; IG_LIKE; 2.
KW   Receptor; Signal.
FT   SIGNAL       1     20       Potential.
FT   CHAIN       21    438       fibroblast growth factor receptor 5
FT                               gamma.
SQ   SEQUENCE   438 AA;  46980 MW;  768D7E96126DD8AE CRC64;

  Query Match             60.3%;  Score 1035.5;  DB 2;  Length 438;
  Best Local Similarity   65.1%;  Pred. No. 7.7e-72;
  Matches  211;  Conservative    4;  Mismatches   14;  Indels   95;  Gaps    2;

Qy          1 MTPSPLLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL 60
              || || |||    ||||| | | |||
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Db          1 MTRSPALLL----LLLGALPSAEAAR-------------------------------- 22

Qy         61 TMWTKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVLDDI 120
                                                                    |||
Db         23 ------------------------------------------------------DDI 25

Qy        121 SPGKESLGPDSSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 180
              ||||||  ||  ||||||||||||||||||||||||||||||||||||||||||||||||
Db         26 SPGKESPGPGGSSGGQEDPASQQWARPRFTQPSKMRRRVIARPVGSSVRLKCVASGHPRP 85

Qy        181 DITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQ 240
              || ||||||  ||  ||:| ||||||||||||:|||||||||||||:|||||||||||||
Db         86 DIMWMKDDQTLTHLEASEHRKKKWTLSLKNLKPEDSGKYTCRVSNKAGAINATYKVDVIQ 145

Qy        241 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGAEGRHNSTIDVGG 300
              ||||||||||||||||||||||||||||||||||||||||||:||||||||||||||||||
Db        146 RTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDVKPVIQWLKRVEYGSEGRHNSTIDVGG 205

Qy        301 QKFVVLPTGDVWSRPDGSYLNKPL 324
              |||||||||||||||||||| |||
Db        206 QKFVVLPTGDVWSRPDGSYLNKLL 229
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RESULT 14
Q6PJN1
ID Q6PJN1 PRELIMINARY; PRT; 350 AA.
AC Q6PJN1;
DT 05-JUL-2004 (TrEMBLrel. 27, Created)
DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
DE FGFRL1 protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Kidney;
RX MEDLINE=22388257; PubMed=12477932;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Kidney;
RA Strausberg R.;
RL Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
DR EMBL; BC013955; AAH13955.1; -.
DR InterPro; IPR003599; Ig.
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 2.
DR SMART; SM00409; IG; 2.
DR SMART; SM00408; IGc2; 2.
DR PROSITE; PS50835; IG_LIKE; 2.
SQ SEQUENCE 350 AA; 37962 MW; C92A18DB4374A831 CRC64;

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  Query Match             52.1%;  Score 894;  DB 2;  Length 350;
  Best Local Similarity   99.4%;  Pred. No. 5.3e-61;
  Matches  169;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy        155 MRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPE 214
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPE 60

Qy        215 DSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDV 274
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 DSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDV 120

Qy        275 KPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        121 KPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKLL 170
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RESULT 15
AAH13955
ID AAH13955 PRELIMINARY; PRT; 350 AA.
AC AAH13955;
DT 02-MAR-2004 (TrEMBLrel. 27, Created)
DT 02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
DT 02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
DE FGFRL1 protein.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primata; Catarrhini; Hominidae; Homo.
OX NCBI_TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=Kidney;
RX MEDLINE=22388257; PubMed=12477932;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Jones S.J., Marra M.A.;
RT "Generation and initial analysis of more than 15,000 full-length human
RT and mouse cDNA sequences.";
RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
RN [2]
RP SEQUENCE FROM N.A.
RC TISSUE=Kidney;
RA Strausberg R.;
RL Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases.
DR EMBL; BC013955; AAH13955.1; -.
SQ SEQUENCE 350 AA; 37962 MW; C92A18DB4374A831 CRC64;

```
  Query Match             52.1%;  Score 894;  DB 2;  Length 350;
  Best Local Similarity   99.4%;  Pred. No. 5.3e-61;
  Matches  169;  Conservative    0;  Mismatches    1;  Indels    0;  Gaps    0;

Qy        155 MRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPE 214
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 MRRRVIARPVGSSVRLKCVASGHPRPDITWMKDDQALTRPEAAEPRKKKWTLSLKNLRPE 60

Qy        215 DSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDV 274
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 DSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVLTGTHPVNTTVDFGGTTSFQCKVRSDV 120
```

```
Qy        275 KPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKPL 324
              ||||||||||||||||||||||||||||||||||||||||||||||||| |
Db        121 KPVIQWLKRVEYGAEGRHNSTIDVGGQKFVVLPTGDVWSRPDGSYLNKLL 170
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Search completed: February 2, 2005, 18:26:02
Job time : 197 secs